

Supplemental File S2 – Heme Biosynthesis

Synthesis of heme is critical to hemoglobin formation and oxygen transport. In a normal man, 6.3 g of hemoglobin is replaced daily with little wastage; porphyrias result from disruption of synthesis or feedbacks in process control, often due to deficiency of one or more enzymes in the synthesis pathway (Rimington 1989). Defects in each of the eight enzymes in the heme biosynthetic pathway have been associated with a specific porphyria (Phillips 2019). We examined variation at nine genes (O’Malley *et al.* 2018) among sciurids.

The *ALAD* gene encodes aminolevulinate dehydratase. Diseases associated with *ALAD* include porphyria, and acute hepatic porphyria. Among its related pathways are porphyrin and innate immune system response. The *ALAD* enzyme is composed of eight identical subunits and catalyzes the condensation of two molecules of δ -aminolevulinate to form porphobilinogen (a precursor of heme). *ALAD* catalyzes the second step in the porphyrin and heme biosynthetic pathway. *ALAD* enzymatic activity is inhibited by lead and a defect in the *ALAD* structural gene can cause increased sensitivity to lead poisoning and acute hepatic porphyria. Alternative splicing of this gene results in multiple transcript variants encoding different isoforms.

Our sequence alignment included 355 predicted amino acids for *S. niger*. There were differences at 16 amino acids among *Sciurus* species; additionally, *S. vulgaris* lacked 23 amino acids at positions 270-293. Three ground squirrels – *S. dauricus*, *I. tridecemlineatus*, and *U. parryii* - had a 32-amino acid sequence at positions 140-172 not seen in other species.

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Alad                MHHQSVLHSGYFHPLLRWQTAASTVVSASNLIYPIFVTDVDPDDVQPIASLPGVARYGVNQ
Aplodontia_rufa    MQPQSILHSGYFHPLLRWQTAATTLASASNLIYPIFVT-VSKN-GPDLAQGGGRCGTGQ
Xerus_inauris@A    MQPQSVLHSGYFHPLLRWQAAATSLASASNFIYPIFVT-VPRNVQPIASLPGVARYGVNQ
Marmota_monax@A    MQPQSVLHSGYFHPLLRWQTAATTLASASNFIYPIFVT-VPDDPASQSLSPPRYGVNR
Spermophilus_da    MQPQSVLHSGYFHPLLRWQTAATTLASASNFIYPIFVT-VPRN-GPDLSVHGGGRYGVNQ
Ictidomys_tride    MQPQSVLHSGYFHPLLRWQTAATTLASASNFIYPIFVT-VPDDVQPPQSLSPPRYGVNR
Urocitellus_par    MQPQSVLHSGYFHPLLRWQTAATTLASASNFIYPIFVT-VPRNVQPPQSLSPPRYGVNR
Sciurus_vulgari    MQPQSVLHSGYFHPLLRWQTAATTLASASNFIYPIFVTDVDPDDIQPIASLPVMARYRVNR
Sciurus_carolin    MQPQSVLHSGYFHPLLRWQTAATTLASASNFIYPIFVTDVDPDDIQPIASLPGMSRYGVNQ
Sciurus_niger@A    MHPQSVLHSGYFHPLLRWQTAATTLASASNFIYPIFVTDVDPDDIQPIANLPGMSRYGVNQ
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Alad                LEEMLRPLVEAGLRCVLI FGVP SRVPKDEQGSAA DSEDSPTIEAVRLLRKT FPSLLVACD
Aplodontia_rufa    LEEMLRPLVEAGLRCVLI FGVP SRVPKDERGSAADSEESPAIEAIRLLRKT FPNLLVACD
Xerus_inauris@A    LEEMLRPLVEAGLRCVLI FGVP SRVP- DERGSAADSEESPAIEAIRLLRKT FPSLLVACD
Marmota_monax@A    LEEMLKPLVEEGLRCVLI FGVP SRVPKDEQGSAA DSEESPTIEAVRLLRKT FPSLLVACD
Spermophilus_da    LEEMLRPLVEAGLRCVLI FGVP SRCP- DERGSAADSEESPTIEAVRLLRKT FPSLLVACD
Ictidomys_tride    LEEMLKPLVEEGLRCVLI FGVP SRVPKDEQGSAA DSEESPTIEAVRLLRKT FPSLLVACD
Urocitellus_par    LEEMLKPLVEEGLRCVLI FGVP SRVPKDEQGSAA DSEESPTIEAVRLLRKT FPSLLVACD
Sciurus_vulgari    LEEMLRPLVEEGLCCVLI FGVP SKVPKDERGSAADSEKSPAIEAIRLLRKT FPNLLVACD
Sciurus_carolin    LEEMLRPLVEAGLRCVLI FGVP SRVPKDERGSAADSEKSPAIEAIHLLRKT FPNLLVACD
Sciurus_niger@A    LEEMLRPLVEAGLRCVLI FGVP SRVPKDERGSAADSEDSPTIEAVRLLRKT FPNLLVACD
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Alad                VCLCPYTSHGHCGLLENGAFLAEESRQRLAEVALAYAKA-----
Aplodontia_rufa    VCLCPYTSHGHCGLLENGAFRAEESRQRLAEVALAYAKA-----
Xerus_inauris@A    VCLCPYTSHGHCGLLENGAFRAEESRQRLAEVALAYAKA-----
Marmota_monax@A    VCLCPYTSHGHCGLLENGAFKAEESRQRLAEVALAYAKA-----
    
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Spermophilus_da VCLCPYTSHGHCGLLENGAFKAEESRQRLAEVALAYAKAGEVPRRGHGLWVWVLAEWIE
 Ictidomys_tride VCLCPYTSHGHCGLLENGAFKAEESRQRLAEVALAYAKAGEVPSRGGHGLWVWVLAEWIE
 Urocitellus_par VCLCPYTSHGHCGLLENGAFKAEESRQRLAEVALAYAKAGEVPSRGGHGLWVWVLAEWIE
 Sciurus_vulgari ICLCPYTSHGHCGLLENGAFRAEESRQRLAEVALAYAKA-----
 Sciurus_carolin ICLCPYTSHGHCGLLENGAFRAEESRQRLAEVALAYAKA-----
 Sciurus_niger@A ICLCPYTSHGHCGLLENGAFRAEESRQRLAEVALAYAKA-----
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Alad -----GCQVVAPSDMMDGRVEAIKAALLKHGLGNRVSVMSYSAKFASCFYGGPF
 Aplodontia_rufa -----GAQVVAPSDMMDGRVEAIKAALLKHGLGNRVSVMSYSAKFASCFYGGPF
 Xerus_inauris@A -----GCQVVAPSDMMDGRVEAIKEALMAHGLGNRVSVMSYSAKFASCFYGGPF
 Marmota_monax@A -----GCQVVAPSDMMDGRVEAIKEALMAHGLGNRVSVMSYSAKFASCFYGGPF
 Spermophilus_da GPRVLKTTLCPSGCQVVAPSDMMDGRVEAIKDALMAHGLGNRVSVMSYSAKFASCFYGGPF
 Ictidomys_tride GPRVLKTTLCPSGCQVVAPSDMMDGRVEAIKDALMAHGLGNRVSVMSYSAKFASCFYGGPF
 Urocitellus_par GPRVLKTTLCPSGCQVVAPSDMMDGRVEAIKDALLAHGLGNRVSVMSYSAKFASCFYGGPF
 Sciurus_vulgari -----GCQVVAPSDMMDGRVEAIKEALMAHGLGNRVSVMSYSAKFASCFYGGPF
 Sciurus_carolin -----GCQVVAPSDMMDGRVEAIKEALMAHGLGNRVSVMSYSAKFASCFYGGPF
 Sciurus_niger@A -----GCQVVAPSDMMDGRVEAIKEALMAHGLGNRVSVMSYSAKFASCFYGGPF
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Alad RDAAQSSPAFGDRRCYQLPPGARGLALRAVA-----
 Aplodontia_rufa RDAAQSSPAFGDRRCYQLPPGARGLALRAVA-----
 Xerus_inauris@A RDAAQSSPAFGDRRCYQLPPGARGLALRAVVDQAA-----THTHPLPA----YPL
 Marmota_monax@A RDAAQSSPAFGDRRCYQLPPGARGLALRAVA-----
 Spermophilus_da RDAAQSSPAFGDRRCYQLPPGARGLALRAVVDQAD-----THTHTHTLPA----CPF
 Ictidomys_tride RDAAQSSPAFGDRRCYQLPPGARGLALRAVVDQAADTHTHTYHTHTHTLPA----CPF
 Urocitellus_par RDAAQSSPAFGDRRCYQLPPGARGLALRAVA-----
 Sciurus_vulgari QDAALSSPTFGDSCCYQLPPGARGLALQAV-----
 Sciurus_carolin RDAAQSSPAFGDRRCYQLPPGARGLALRAVVDQAA-----THSPCLSPSRINAV
 Sciurus_niger@A RDAAQSSPAFGDRRCYQLPPGARGLALRAVVDQAA-----THSPCLSPSRINAV
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Alad -----RDIQEGADMLMVKPLPYLDMV-REVKDKHPPLPLAVYQVSGEFAMLWHGA
 Aplodontia_rufa -----RDIQEGADMLMVKPLPYLDMV-REVKDQHPPLPLAVYHVSGEYAMLWHGA
 Xerus_inauris@A PRHCTLIIPDRDVREGADMLMVKPGMPYLDIV-REVKDKHPPLPLAVYHVSGEFAMLWHGA
 Marmota_monax@A -----RDIQEGADMLMVKPLPYLDMV-REVKDQHPQLPLAVYHVSGEFAMLWHGA
 Spermophilus_da PRHFTLFPDRDVREGADMLMVKPLPYLDMV-REVKDKHPQLPLAVYHVSGEFAMLWHGA
 Ictidomys_tride PRHFTLFPDRDVREGADMLMVKPGMPYLDIV-REVKDKHPQLPLAVYHVSGEFAMLWHGA
 Urocitellus_par -----RDIQEGADMLMVKPLPHLNPVMAASPQHPQLPLAVYHVSGEFAMLWHGA
 Sciurus_vulgari -----DRDVQEGADMLMVKPGMPYLDIV-WEVKDKHPPLPLAVYHVSGEFAMLWHGA
 Sciurus_carolin VRHFTFIIPDRDVQEGADMLMVKPGMPYLDIV-REVKDKHPPLPLAVYHVSGEFAMLWHGA
 Sciurus_niger@A VRHFTLIIPDRDVQEGADMLMVKPGMPYLDIV-REVKDKHPPLPLAVYHVSGEFAMLWHGA
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Alad QAGAFDLRTAVLETMTAFRRAGADIIITYFAPQLLKWKEEZ
 Aplodontia_rufa RAGAFDLKAAVLEAMTAFRRAG-----
 Xerus_inauris@A RAGAFDLKAAVLEAMTAFRRAG-----
 Marmota_monax@A RAGAFDLKAAVLEAMTAFRRAG-----
 Spermophilus_da RAGAFDLKAAVLEAMTAFRRAG-----
 Ictidomys_tride RAGAFDLKAAVLEAMTAFRRAG-----
 Urocitellus_par RAGAFDLKAAVLEAMTAFRRAG-----
 Sciurus_vulgari QAGAFDLKAAVLEAMTAFRRAGADVIITYHTPQLLQWLKE--
 Sciurus_carolin QAGAFDLKAAVLEAMTAFRRAGADIIITYTPQLLQWLKE--
 Sciurus_niger@A QAGAFDLKAAVLEAMTAFRRAGADIIITYTPQLLRWLKE--
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ALAS1 encodes 5'-aminolevulinatase synthase 1. Diseases associated with *ALAS1* include acute porphyria and anemia, sideroblastic, 1. Among its related pathways are porphyrin and glycine, serine, and threonine metabolism. Gene ontology (GO) annotations related to this gene include pyridoxal phosphate binding and 5-aminolevulinatase synthase activity. An important paralog of this gene is *ALAS2*. This gene encodes the mitochondrial enzyme which catalyzes the rate-limiting step in heme (iron-protoporphyrin) biosynthesis. The enzyme encoded by this gene is the housekeeping enzyme; a separate gene encodes a form of the enzyme that is specific for erythroid tissue. The level of the mature encoded protein is regulated by heme; high levels of heme down-regulate the mature enzyme in mitochondria, while low heme levels up-regulate it. Alternative splicing results in multiple transcript variants encoding different isoforms.

The *ALAS1* amino acid sequence was rather highly conserved among sciurids and among all species considered. Among 640 amino acids for *Sciurus* species, we observed three point differences. The human sequence had a two-amino acid insertion at sites 84-85.

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Alas1      METVVRRCPFLLSRVPQAFLLQKAGKSLFFYAQNCPKMMVEVGAKPAPRTLSTSAVHCQQVKE
Aplodontia_rufa METVVRRCPFLLSRVPQAFLLQKAGKSLFFYAQNCPKMMVEVWSPVPRALTTSAVHCQQIIE
Xerus_inauris@A METVVRRCPFLLSRVPQAFLLQKAGKSLFFYAQNCPKMMIEIGSKPAPRALSTSAVHCQQIIE
Marmota_monax@A METVVRRCPFLLSRVPQAFLLQKAGKSLFFYAQNCPKMMVEVGSKPVPRAVSTSAVHCQQIIE
Spermophilus_da METVVRRCPFLLSRVPQAFLLQKAGKSLFFYAQNCPKMMVEVGSKPVPRAVSTSAVHCQQIIE
Ictidomys_tride METVVRRCPFLLSRVPQAFLLQKAGKSLFFYAQNCPKMMVEVGSKPVPRAVSTSAVHCQQIIE
Urocitellus_par METVVRRCPFLLSRVPQAFLLQKAGKSLFFYAQNCPKMMVEVGSKPVPRAVSTSAVHCQQIIE
Sciurus_vulgari METVVRRCPFLLSRVPQAFLLQKAGKSLFFYAQNCPKMMVEVGSKPAPRALSTSAVHCQQIIE
Sciurus_carolin METVVRRCPFLLSRVPQAFLLQKAGKSLFFYAQNCPKMMVEVGSKPAPRALSTSAVHCQQIIE
Sciurus_niger@A METVVRRCPFLLSRVPQAFLLQKAGKSLFFYAQNCPKMMVEVGSKPAPRALSTSAVHCQQIIE
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Alas1      TPPANEKEKTAKAAVQQAPDESQMAQTPDGTQLPSGHSPATSQGSGSKCPFLAAQLSQT
Aplodontia_rufa TPPAGEEDKTAKAKVQETPDRSQ--QTPNGTQLPSAHLSPAMSQGTSSKCPFLAAQMSQT
Xerus_inauris@A TPPASEEDKTAKTKVQQAPDRSQ--QTPNGMQLPSGHPLPAMSQGTASKCPFLAAQMSQT
Marmota_monax@A TPPASEEDKTAKAKVQQAPDRSQ--QTPNGTQLPSGHHS PAMSQGTASKCPFLAAQMSQT
Spermophilus_da TPPASEEDKTAKAKVQQAPDRSQ--QTPNGTQLPSGHHS PAMSQGTASKCPFLAAQMSQT
Ictidomys_tride TPPASEEDKTAKAKVQQAPDRSQ--QTPNGTQLPPGHHS PAMSQGTASKCPFLAAQMSQT
Urocitellus_par TPPASEEDKTAKAKVQQAPDRSQ--QTPNGTQLPSGHHS LAMSQGTASKCPFLAAQMSQT
Sciurus_vulgari TPPASEDKTANKSKVQQAPDRSQ--QTPNGTQLPSGHSPAMSQGTASKCPFLAAQMSQT
Sciurus_carolin TPPASEDKTANKSKVQQAPDRSQ--QTPNGTQLPSGHSPAMSQGTASKCPFLAAQMSQT
Sciurus_niger@A TPPASEDKTANKSKVQQAPDRSQ--QTPNGTQLPSGHSPAVSQGTASKCPFLAAQMSQT
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Alas1      GSSVFRKASLELQEDVQEMHAVRKEAAQSPVPPSLVNVKTDGEDPSRLLKNFQDIMRKQR
Aplodontia_rufa GSSVFCASLELQEDVQEMHAVRKEVAQTSVKPTVINVKTNDDPSGLLKNFQDIMRKQR
Xerus_inauris@A GSNVFCASLELQEDVQEMHAVRKEVAQTSVNP SVI SVKTDGKDPSELLKNFQDIMQKQR
Marmota_monax@A GSNVFCASLELQEDVQEMHAVRKEVAQTSVNP NV INVKTDGEDPSELLKNFQDIMRKQR
Spermophilus_da GSNVFCASLELQEDVQEMHAVRKEVAQTSVNP NV INVKTDGEDPSELLKNFQDIMRKQR
Ictidomys_tride GSNVFCASLELQEDVQEMHAVRKEVAQTSVNP NV INVKTDGEDPSELLKNFQDIMRKQR
Urocitellus_par GSNVFCASLELQEDVQEMHAVRKEVAQTSVNP NV INVKTDGEDPSELLKNFQDIMRKQR
Sciurus_vulgari GSSVFCASLELQEDVQEMHAVRKEVAQTSVNP NVI SVKTDGEDPNELLKNFQDIMRKQR
Sciurus_carolin GSSVFCASLELQEDVQEMHAVRKEVAQTSVNP SVI SVKTDGEDPNELLKNFQDIMRKQR
Sciurus_niger@A GSSVFCASLELQEDVQEMHAVRKEVAQTSVNP NVI SVKTDGEDPNELLKNFQDIMRKQR
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Alas1      PERVSHLLQDNLPKSVSTFQYDHF FEKKI DEKKN DHTYRVFKTVNRR AQIFPMADDY TDS
Aplodontia_rufa PERVSHLLQDNLPKSVSTFQYDHF FEKKI DEKKN DHTYRVFKTVNRR AQIFPMADDY SDS
Xerus_inauris@A PERVSHLLQDNLPKSVSTFQYDHF FEKKI DEKKN DHTYRVFKTVNRR AQIFPMADDY SDS
    
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Marmota_monax@A PERVSHLLQDNLPKSVSTFQYDHFPEKKIDEKKNNDHTYRVFKTVNRRQIFPMADDYSDS
Spermophilus_da PERVSHLFQDNLPKSVSTFQYDHFPEKKIDEKKNNDHTYRVFKTVNRRQIFPMADDYSDS
Ictidomys_tride PERVSHLLQDNLPKSVSTFQYDHFPEKKIDEKKNNDHTYRVFKTVNRRQIFPMADDYSDS
Urocitellus_par PERVSHLLQDNLPKSVSTFQYDHFPEKKIDEKKNNDHTYRVFKTVNRRQIFPMADDYSDS
Sciurus_vulgari PERVSHLLQDNLPKSVSTFQYDHFPEKKIDEKKNNDHTYRVFKTVNRRQIFPMADDYSDS
Sciurus_carolin PERVSHLLQDNLPKSVSTFQYDHFPEKKIDEKKNNDHTYRVFKTVNRRQIFPMADDYSDS
Sciurus_niger@A PERVSHLLQDNLPKSVSTFQYDHFPEKKIDEKKNNDHTYRVFKTVNRRQIFPMADDYSDS
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Alas1 LITKKQVSVWCSNDYLGMRSRHRVCGAVMETVKQHGGAGGTRNISGTSKHFHVELEQALA
Aplodontia_rufa LITKKQVSVWCSNDYLGMRSRHRVCGAVMETVKQHGGAGGTRNISGTSKHFHVELEQELA
Xerus_inauris@A LITKKQVSVWCSNDYLGMRSRHRVCGAVMETVKQHGGAGGTRNISGTSKHFHVELEQELA
Marmota_monax@A LITKKQVSVWCSNDYLGMRSRHRVCGAVMETVKQHGGAGGTRNISGTSKHFHVELEQELA
Spermophilus_da LITKKQVSVWCSNDYLGMRSRHRVCGAVVETLKHGGAGGTRNISGTSKHFHVELEQELA
Ictidomys_tride LITKKQVSVWCSNDYLGMRSRHRVCGAVMETVKQHGGAGGTRNISGTSKHFHVELEQELA
Urocitellus_par LITKKQVSVWCSNDYLGMRSRHRVCGAVMETVKQHGGAGGTRNISGTSKHFHVELEQELA
Sciurus_vulgari LITKKQVSVWCSNDYLGMRSRHRVCGAVMETVKQHGGAGGTRNISGTSKHFHVELEQELA
Sciurus_carolin LITKKQVSVWCSNDYLGMRSRHRVCGAVMETVKQHGGAGGTRNISGTSKHFHVELEQELA
Sciurus_niger@A LITKKQVSVWCSNDYLGMRSRHRVCGAVMETVKQHGGAGGTRNISGTSKHFHVELEQELA
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Alas1 DLHGKDAALLFSSCFVANDSTLFTLAKMMPGCEIYSDSGNHASMIQGIRNSRVPKYIFRHR
Aplodontia_rufa DLHGKDAALLFSSCFVANDSTLFTLAKMMPGCEIYSDSGNHASMIQGIRNSRVPKYIFRHR
Xerus_inauris@A DLHGKDAALLFSSCFVANDSTLFTLAKMMPGCEIYSDSGNHASMIQGIRNSRVPKYIFRHR
Marmota_monax@A DLHGKDAALLFSSCFVANDSTLFTLAKMMPGCEIYSDSGNHASMIQGIRNSRVPKYIFRHR
Spermophilus_da DLHGKDAALLFSSCFVANDSTLFTLAKMMPGCEIYSDSGNHASMIQGIRNSRVPKYIFRHR
Ictidomys_tride DLHGKDAALLFSSCFVANDSTLFTLAKMMPGCEIYSDSGNHASMIQGIRNSRVPKYIFRHR
Urocitellus_par DLHGKDAALLFSSCFVANDSTLFTLAKMMPGCEIYSDSGNHASMIQGIRNSRVPKYIFRHR
Sciurus_vulgari DLHGKDAALLFSSCFVANDSTLFTLAKMMPGCEIYSDSGNHASMIQGIRNSRVPKYIFRHR
Sciurus_carolin DLHGKDAALLFSSCFVANDSTLFTLAKMMPGCEIYSDSGNHASMIQGIRNSRVPKYIFRHR
Sciurus_niger@A DLHGKDAALLFSSCFVANDSTLFTLAKMMPGCEIYSDSGNHASMIQGIRNSRVPKYIFRHR
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Alas1 NDVNHLRELLQRSDPVSPKIVAFETVHSMGDGAVCPLEELCDVAHEFGAIFVDEVHAVGL
Aplodontia_rufa NDVNHLRELLQRSDPVSPKIVAFETVHSMGDGAVCPLEELCDVAHEFGAIFVDEVHAVGL
Xerus_inauris@A NDVNHLRELLQRSDPVSPKIVAFETVHSMGDGAVCPLEELCDVAHEFGAIFVDEVHAVGL
Marmota_monax@A NDVNHLRELLQRSDPVSPKIVAFETVHSMGDGAVCPLEELCDVAHEFGAIFVDEVHAVGL
Spermophilus_da NDVNHLRELLQRSDPVSPKIVAFETVHSMGDGAVCPLEELCDVAHEFGAIFVDEVHAVGL
Ictidomys_tride NDVNHLRELLQRSDPVSPKIVAFETVHSMGDGAVCPLEELCDVAHEFGAIFVDEVHAVGL
Urocitellus_par NDVNHLRELLQRSDPVSPKIVAFETVHSMGDGAVCPLEELCDVAHEFGAIFVDEVHAVGL
Sciurus_vulgari NDVNHLRELLQRSDPVSPKIVAFETVHSMGDGAVCPLEELCDVAHEFGAIFVDEVHAVGL
Sciurus_carolin NDVNHLRELLQRSDPVSPKIVAFETVHSMGDGAVCPLEELCDVAHEFGAIFVDEVHAVGL
Sciurus_niger@A NDVNHLRELLQRSDPVSPKIVAFETVHSMGDGAVCPLEELCDVAHEFGAIFVDEVHAVGL
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Alas1 YGARGGGIGDRDGVMPKMDIISGTLGKAFGCGVGGYIASTSLIDIVRSYAAGFI FT TSLP
Aplodontia_rufa YGARGGGIGDRDGVMPKMDIISGTLGKAFGCGVGGYIASTSLLDIVRSYAAGFI FT TSLP
Xerus_inauris@A YGAQGGGIGDRDGVMPKMDIISGTLGKAFGCGVGGYIASTSLIDIVRSYAAGFI FT TSLP
Marmota_monax@A YGARGGGIGDRDGVMPKMDIISGTLGKAFGCGVGGYIASTSLIDIVRSYAAGFI FT TSLP
Spermophilus_da YGARGGGIGDRDGVMPKMDIISGTLGKAFGCGVGGYIASTSLIDIVRSYAAGFI FT TSLP
Ictidomys_tride YGARGGGIGDRDGVMPKMDIISGTLGKAFGCGVGGYIASTSLIDIVRSYAAGFI FT TSLP
Urocitellus_par YGARGGGIGDRDGVMPKMDIISGTLGKAFGCGVGGYIASTSLIDIVRSYAAGFI FT TSLP
Sciurus_vulgari YGAQGGGIGDRDGVMPKMDIISGTLGKAFGCGVGGYIASTSLIDIVRSYAAGFI FT TSLP
Sciurus_carolin YGAQGGGIGDRDGVMPKMDIISGTLGKAFGCGVGGYIASTSLIDIVRSYAAGFI FT TSLP
Sciurus_niger@A YGAQGGGIGDRDGVMPKMDIISGTLGKAFGCGVGGYIASTSLIDIVRSYAAGFI FT TSLP
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Alas1          PMLLAGALESVRILKSSEGRALRRQHQRNVKLLRQMLMDAGLPVIHCPSHIIPVRVADAA
Aplodontia_rufa PMLLAGALESVRILKSSEGRALRRQHQRNVKLMRQMLMDAGLPVIHCPSHIIPVRVADAA
Xerus_inauris@A PMLLAGALESVRILKSSEGRALRRQHQRNVKLMRQMLMDAGLPVIHCPSHIIPVRVADAA
Marmota_monax@A PMLLAGALESVRILKSSEGRALRRQHQRNVKLMRQMLMDAGLPVIHCPSHIIPVRVADAA
Spermophilus_da PMLLAGALESVRILKSSEGRALRRQHQRNVKLMRQMLMDAGLPVIHCPSHIIPVRVADAA
Ictidomys_tride PMLLAGALESVRILKSSEGRALRRQHQRNVKLMRQMLMDAGLPVIHCPSHIIPVRVADAA
Urocitellus_par PMLLAGALESVRILKSSEGRALRRQHQRNVKLMRQMLMDAGLPVIHCPSHIIPVRVADAA
Sciurus_vulgari PMLLAGALESVRILKSSEGRALRRQHQRNVKLMRQMLMDAGLPVIHCPSHIIPVRVADAA
Sciurus_carolin PMLLAGALESVRILKSSEGRALRRQHQRNVKLMRQMLMDAGLPVIHCPSHIIPVRVADAA
Sciurus_niger@A PMLLAGALESVRILKSSEGRALRRQHQRNVKLMRQMLMDAGLPVIHCPSHIIPVRVADAA
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Alas1          KNTEICDELMTRHNIYVQAINYPTVPRGEELLRIAPTPHHTPQMMNFFVEKLLVTWKRVG
Aplodontia_rufa KNTEICDELMTRHNIYVQAINYPTVPRGEELLRIAPTPHHTPQMMNFFVEKLLVTWKRVG
Xerus_inauris@A KNTEICDELMTRHNIYVQAINYPTVPRGEELLRIAPTPHHTPQMMNFFVEKLLVTWKRVG
Marmota_monax@A KNTEICDELMTRHNIYVQAINYPTVPRGEELLRIAPTPHHTPQMMNFFVEKLLVTWKRVG
Spermophilus_da KNTEICDELMTRHNIYVQAINYPTVPRGEELLRIAPTPHHTPQMMNFFVEKLLVTWKRVG
Ictidomys_tride KNTEICDELMTRHNIYVQAINYPTVPRGEELLRIAPTPHHTPQMMNFFVEKLLVTWKRVG
Urocitellus_par KNTEICDELMTRHNIYVQAINYPTVPRGEELLRIAPTPHHTPQMMNFFVEKLLVTWKRVG
Sciurus_vulgari KNTEICDELMTRHNIYVQAINYPTVPRGEELLRIAPTPHHTPQMMNFFVEKLLVTWKRVG
Sciurus_carolin KNTEICDELMTRHNIYVQAINYPTVPRGEELLRIAPTPHHTPQMMNFFVEKLLVTWKRVG
Sciurus_niger@A KNTEICDELMTRHNIYVQAINYPTVPRGEELLRIAPTPHHTPQMMNFFVEKLLVTWKRVG
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Alas1          LELKPHSSAECNFCRRPLHFEVMSEREKAYFSGMSKLVSAQAZ
Aplodontia_rufa LELKPHSSAECNFCRRPLHFEVMSEREKAYFSGMSKLVSAQA-
Xerus_inauris@A LELKPHSSAECNFCRRPLHFEVMSEREKAYFSGMSKLVSAQA-
Marmota_monax@A LELKPHSSAECNFCRRPLHFEVMSEREKAYFSGMSKLVSAQA-
Spermophilus_da LELKPHSSAECNFCRRPLHFEVMSEREKAYFSGMSKLVSAQA-
Ictidomys_tride LELKPHSSAECNFCRRPLHFEVMSEREKAYFSGMSKLVSAQA-
Urocitellus_par LELKPHSSAECNFCRRPLHFEVMSEREKAYFSGMSKLVSAQA-
Sciurus_vulgari LELKPHSSAECNFCRRPLHFEVMSEREKAYFSGMSKLVSAQA-
Sciurus_carolin LELKPHSSAECNFCRRPLHFEVMSEREKAYFSGMSKLVSAQA-
Sciurus_niger@A LELKPHSSAECNFCRRPLHFEVMSEREKAYFSGMSKLVSAQA-
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ALAS2 encodes 5'-aminolevulinic synthase 2. Diseases associated with *ALAS2* include anemia, sideroblastic, 1 and protoporphyria, erythropoietic, X-linked pyridoxine-responsive sideroblastic anemia. Among its related pathways are porphyrin and glycine, serine, and threonine metabolism. Gene ontology (GO) annotations related to this gene include pyridoxal phosphate binding and glycine binding. An important paralog of this gene is *ALAS1*. The product of this gene specifies an erythroid-specific mitochondrially located enzyme. The encoded protein catalyzes the first step in the heme biosynthetic pathway. Alternatively spliced transcript variants encoding different isoforms have been identified.

Among 586 amino acids, we observed three differences among *Sciurus* species at *ALAS2*. We noted an insertion of 15-16 amino acids at positions 61-76 in seven non-*Sciurus* sequences. *X. inauris* had an insertion of five amino acids at positions 554-559.

```

Alas2          MVAAAMLLRSCPVLQSGPTGLLGKVKAKTYQFLFSIGRCPILATQGPICSIHLKATKAGG
Aplodontia_rufa MVAAAVLLQCCPVLARGPTGLLGKVIKTHQFLFGTGRCPIAQAQGPICSIHLKATKAGG
Xerus_inauris@A MVAAAMLLQCCPVLARGPTGLLGKVIKTHQFLFGTGRCPIAQAQGPICSIHLKTTNAGG
Marmota_monax@A MVAAAVLLQCCPVLARGPTGLLSKVVKTHQFLFGTGRCPIAQAQGPICSIHLKATRAGG

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Spermophilus_da MVAAAVLLQCCPVLARGPTGLLSKVVKTHQFLFGTGRCPI LATQGPICSQIHLKATRAGG
Ictidomys_tride MVAAAVLLQCCPVLARGPTGLLSKVIKTHQFLFGTGRCPI LATQGPICSQIHLKATRAGG
Urocitellus_par MVAAAVLLQCCPVLARGPTGLLSKVVKTHQFLFGTGRCPI LATQGPICSQIHLKATRAGG
Sciurus_vulgari MVAAAVLLQCCPVLARGPTGLLGKVVKTHQFLFGTGRCPI LAAQGPICSQIHLKATKAGG
Sciurus_carolin MVAAAVLLQCCPVLARGPTGLLGKVVKTHQFLFGTGRCPI LATQGPICSQIHLKATKAGG
Sciurus_niger@A MVAAAVLLQCCPVLVRGPTVLLGKVVKTHQFLFGTGRCPI LATQGPICSQIHLKATKAGG
*****:***:**** :*** **.* ** :****. *****:*** *****:*.***

Alas2 -----DSPSWAKSHCPFMLSELQDRKSKIVQRAAPEVQEDVKTFKTDLL
Aplodontia_rufa -----DSPSWAKGHCPFMLSELQDGKSKIVQKAAPEVQEDVKTFKTDLL
Xerus_inauris@A -----DSPSWTKGHCPFMLSELQDGKSKIVQKAAPEVQEDVKTFKTDLL
Marmota_monax@A GKKRFLEKGGCLGHSDDS DSPSWAKSHCPFMLSELQDRKSKIVQKAAPEVQEDVKTFKTDLL
Spermophilus_da GKKRFLEKGGCLSHS- DSPSWAKSHCPFMLSELQDRKSKIVQKAAPEVQEDVKTFKTDLL
Ictidomys_tride GKKRFLEKGGCLGHS- DSPSWAKSHCPFMLSELQDRKSKIVQKAAPEVQEDVKTFKTDLL
Urocitellus_par GKKRFLEKGGCLGHS- DSPSWAKSHCPFMLSELQDRKSKIVQKAAPEVQEDVKTFKTDLL
Sciurus_vulgari -----DSPSWSKGHCPFMLSELQDGKSKIVQKAAPEVQEDVKTFKTDLL
Sciurus_carolin -----DSPSWSKGHCPFMLSELQDGKSKIVQKAAPEVQEDVKTFKTDLL
Sciurus_niger@A -----DSPSWSKGHCPFMLSELQDGKSKIVQKAAPEVQEDVKTFKTDLL
*****:*.***** *****:*****

Alas2 STMDSTTRSHSFPSFQEPETE GAVPHLIQNNMTGSQAFGYDQFFRDKIMEKKQDHTYRV
Aplodontia_rufa STMDSTTRSHSFPSFQEPETE GAVPHLIQNNMTGEHAFGYDQFFRDKIMEKKQDHTYRV
Xerus_inauris@A STMDSTTRSHSFPSFQEPETE GAVPHLIQNNMTGEHAFGYDQFFRDKIMEKKQDHTYRV
Marmota_monax@A STMDSTTRSHSFPSFQEPETE GAVPHLIQNNMTGEHAFGYDQFFRDKIMEKKQDHTYRV
Spermophilus_da STMDSTTRSHSFPSFQEPETE GAVPHLIQNNMTGEHAFGYDQFFRDKIMEKKQDHTYRV
Ictidomys_tride STMDSTTRSHSFPSFQEPETE GAVPHLIQNNMTGEHAFGYDQFFRDKIMEKKQDHTYRV
Urocitellus_par STMDSTTRSHSFPSFQEPETE GAVPHLIQNNMTGEHAFGYDQFFRDKIMEKKQDHTYRV
Sciurus_vulgari STMDSTTRSHSFPSFQEPETE GAVPHLIQSNPAGEHAFGYDQFFRDKIMEKKQDHTYRV
Sciurus_carolin STMDSTTRSHSFPSFQEPETE GAVPHLIQSNPAGEHAFGYDQFFRDKIMEKKQDHTYRV
Sciurus_niger@A STMDSTTRSHSFPSFQEPETE GAVPHLIQSNPAGEHAFGYDQFFRDKIMEKKQDHTYRV
*****:*.***** *****:*****

Alas2 FKTVNRWANAYPFAQH FSEASMASKDVS VWCSNDYLGISRHRV LQAI EETLKNHGAGAG
Aplodontia_rufa FKTVNRWANAYPFAQH FSEASMASKDVS VWCSNDYLGMSRHRV LQAI QETLKNHGAGAG
Xerus_inauris@A FKTVNRWANAYPFAQH FSEASMASKDVS VWCSNDYLGMSRHRV LQAI QETLKNHGAGAG
Marmota_monax@A FKTVNRWANAYPFAQH FSEASMASKDVS VWCSNDYLGISRHRV LQAI EETLKNHGAGAG
Spermophilus_da FKTVNRWANAYPFAQH FSEASMASKDVS VWCSNDYLGISRHRV LQAI EETLKNHGAGAG
Ictidomys_tride FKTVNRWANAYPFAQH FSEASMASKDVS VWCSNDYLGMSRHRV LQAI QETLKNHGAGAG
Urocitellus_par FKTVNRWANAYPFAQH FSEASMASKDVS VWCSNDYLGMSRHRV LQAI QETLKNHGAGAG
Sciurus_vulgari FKTVNRWANAYPFAQH FSEASMASKDVS VWCSNDYLGISRHRV LQAI EETLKNHGAGAG
Sciurus_carolin FKTVNRWANAYPFAQH FSEASMASKDVS VWCSNDYLGMSRHRV LQAI QETLKNHGAGAG
Sciurus_niger@A FKTVNRWANAYPFAQH FSEASMASKDVS VWCSNDYLGMSRHRV LQAI QETLKNHGAGAG
*****:*****:*****

Alas2 GTRNISGTSKFHVELEQELAE LHQKDSALLFSSCFVANDSTLFTLAKLLPGCEI YSDAGN
Aplodontia_rufa GTRNISGTSKFHVELEQELAE LHQKDSALLFSSCFVANDSTLFTLAKLLPGCEI YSDAGN
Xerus_inauris@A GTRNISGTSKFHVELEQELAE LHQKDSALLFSSCFVANDSTLFTLAKLLPGCEI YSDAGN
Marmota_monax@A GTRNISGTSKFHVELEQELAE LHQKDSALLFSSCFVANDSTLFTLAKLLPGCEI YSDAGN
Spermophilus_da GTRNISGTSKFHVELEQELAE LHQKDSALLFSSCFVANDSTLFTLAKLLPGCEI YSDAGN
Ictidomys_tride GTRNISGTSKFHVELEQELAE LHQKDSALLFSSCFVANDSTLFTLAKLLPGCEI YSDAGN
Urocitellus_par GTRNISGTSKFHVELEQELAE LHQKDSALLFSSCFVANDSTLFTLAKLLPGCEI YSDAGN
Sciurus_vulgari GTRNISGTSKFHVELEQELAE LHQKDSALLFSSCFVANDSTLFTLAKLLPGCEI YSDAGN
Sciurus_carolin GTRNISGTSKFHVELEQELAE LHQKDSALLFSSCFVANDSTLFTLAKLLPGCEI YSDAGN
Sciurus_niger@A GTRNISGTSKFHVELERELAE LHQKDSALLFSSCFVANDSTLFTLAKLLPGCEI YSDAGN
*****:*****:*****

Alas2 HASMIQGIRNSGAAKFVFRHNDPGHLKLLLEKSDPKTPKIVAFETVHSMGDAICPLEELC
Aplodontia_rufa HASMIQGIRNSGAAKFVFRHNDPGHLKLLLEKSDPKTPKIVAFETVHSMGDAICPLEELC
Xerus_inauris@A HASMIQGIRNSGAAKFVFRHNDPGHLKLLLEKSDPKTPKIVAFETVHSMGDAICPLEELC
Marmota_monax@A HASMIQGIRNSGAAKFVFRHNDPGHLKLLLEKSDPKTPKIVAFETVHSMGDAICPLEELC
Spermophilus_da HASMIQGIRNSGAAKFVFRHNDPGHLKLLLEKSDPKTPKIVAFETVHSMGDAICPLEELC
Ictidomys_tride HASMIQGIRNSGAAKFVFRHNDPGHLKLLLEKSDPKTPKIVAFETVHSMGDAICPLEELC
Urocitellus_par HASMIQGIRNSGAAKFVFRHNDPGHLKLLLEKSDPKTPKIVAFETVHSMGDAICPLEELC
Sciurus_vulgari HASMIQGIRNSGAAKFVFRHNDPGHLKLLLEKSDPKTPKIVAFETVHSMGDAICPLEELC
Sciurus_carolin HASMIQGIRNSGAAKFVFRHNDPGHLKLLLEKSDPKTPKIVAFETVHSMGDAICPLEELC
Sciurus_niger@A HASMIQGIRNSGAAKFVFRHNDPGHLKLLLEKSDPKTPKIVAFETVHSMGDAICPLEELC

Alas2 DVAHQYGALTFVDEVHAVGLYGARGAGIGERDGMHKLDIISGTLGKAFGCVGGYIASTR
Aplodontia_rufa DVAHQYGALTFVDEVHAVGLYGARGAGIGERDGMHKLDIISGTLGKAFGCVGGYIASTR
Xerus_inauris@A DVAHQYGALTFVDEVHAVGLYGARGAGIGERDGMHKLDIISGTLGKAFGCVGGYIASTR
Marmota_monax@A DVAHQYGALTFVDEVHAVGLYGARGAGIGERDGMHKLDIISGTLGKAFGCVGGYIASTR
Spermophilus_da DVAHQYGALTFVDEVHAVGLYGARGAGIGERDGMHKLDIISGTLGKAFGCVGGYIASTR
Ictidomys_tride DVAHQYGALTFVDEVHAVGLYGARGAGIGERDGMHKLDIISGTLGKAFGCVGGYIASTR
Urocitellus_par DVAHQYGALTFVDEVHAVGLYGARGAGIGERDGMHKLDIISGTLGKAFGCVGGYIASTR
Sciurus_vulgari DVAHQYGALTFVDEVHAVGLYGARGAGIGERDGMHKLDIISGTLGKAFGCVGGYIASTR
Sciurus_carolin DVAHQYGALTFVDEVHAVGLYGARGAGIGERDGMHKLDIISGTLGKAFGCVGGYIASTR
Sciurus_niger@A DVAHQYGALTFVDEVHAVGLYGARGAGIGERDGMHKLDIISGTLGKAFGCVGGYIASTR

Alas2 DLVDMVRSYAAGFI FTTS LPPMVL SGALESVRLKGEEGQALRAHQNVKHMQRLLMDR
Aplodontia_rufa DLVDMVRSYAAGFI FTTS LPPMVL SGALESVRLKGEEGQALRAHQNVKHMQRLLMDR
Xerus_inauris@A DLVDMVRSYAAGFI FTTS LPPMVL SGALESVRLKGEEGQALRAHQNVKHMQRLLMDR
Marmota_monax@A DLVDMVRSYAAGFI FTTS LPPMVL SGALESVRLKGEEGQALRAHQNVKHMQRLLMDR
Spermophilus_da DLVDMVRSYAAGFI FTTS LPPMVL SGALESVRLKGEEGQALRAHQNVKHMQRLLMDR
Ictidomys_tride DLVDMVRSYAAGFI FTTS LPPMVL SGALESVRLKGEEGQALRAHQNVKHMQRLLMDR
Urocitellus_par DLVDMVRSYAAGFI FTTS LPPMVL SGALESVRLKGEEGQALRAHQNVKHMQRLLMDR
Sciurus_vulgari DLVDMVRSYAAGFI FTTS LPPMVL SGALESVRLKGEEGQALRAHQNVKHMQRLLMDR
Sciurus_carolin DLVDMVRSYAAGFI FTTS LPPMVL SGALESVRLKGEEGQALRAHQNVKHMQRLLMDR
Sciurus_niger@A DLVDMVRSYAAGFI FTTS LPPMVL SGALESVRLKGEEGQALRAHQNVKHMQRLLMDR

Alas2 GFPVIPCPSHI IPIRVGNAALNSKICDLLLLSKHSIYVQAINYPTVPRGEELLRLAPSPHH
Aplodontia_rufa GLPVIIPCPSHI IPIRVGNAALNSKICDLLLLSKHSIYVQAINYPTVPRGEELLRLAPSPHH
Xerus_inauris@A GLPVIIPCPSHI IPIRVGDAALNSKICDLLLLSKHSIYVQAINYPTVPRGEELLRLAPSPHH
Marmota_monax@A GFPVIPCPSHI IPIRVGDAALNSKICDLLLLSKHSIYVQAINYPTVPRGEELLRLAPSPHH
Spermophilus_da GLPVIIPCPSHI IPIRVGDAALNSKICDLLLLSKHSIYVQAINYPTVPRGEELLRLAPSPHH
Ictidomys_tride GFPVIPCPSHI IPIRVGDAALNSKICDLLLLSKHSIYVQAINYPTVPRGEELLRLAPSPHH
Urocitellus_par GLPVIIPCPSHI IPIRVGDAALNSKICDLLLLSKHSIYVQAINYPTVPRGEELLRLAPSPHH
Sciurus_vulgari GLPVIIPCPSHI IPIRVGDAALNSKICDLLLLSKHSIYVQAINYPTVPRGEELLRLAPSPHH
Sciurus_carolin GFPVIPCPSHI IPIRVGDAALNSKICDLLLLSKHSIYVQAINYPTVPRGEELLRLAPSPHH
Sciurus_niger@A GLPVIIPCPSHI IPIRVGDAALNSKICDLLLLSKHSIYVQAINYPTVPRGEELLRLAPSPHH
*:*****:*****

Alas2 SPQMMENFVEKLL-----LAWTEVGLPLQDVSVAACNFCHRPVHFELMSEWERSYFGNMG
Aplodontia_rufa SPQMMENFVEKLL-----LAWTEVGLPLQDVSVAACNFCHRPVHFELMSEWERSYFGNMG
Xerus_inauris@A SPQMMENFVGKYLQSCGSPFGTICKLSLQCLSL-----
Marmota_monax@A SPQMMENFVEKLL-----LAWTEVGLPLQDVSVAACNFCHRPVHFELMSEWERSYFGNMG
Spermophilus_da SPQMMENFVEKLL-----LAWTEVGLPLQDVSVAACNFCHRPVHFELMSEWERSYFGNMG
Ictidomys_tride SPQMMENFVEKLL-----LAWTEVGLPLQDVSVAACNFCHRPVHFELMSEWERSYFGNMG
Urocitellus_par SPQMMENFVEKLL-----LAWTEVGLPLQDVSVAACNFCHRPVHFELMSEWERSYFGNMG
Sciurus_vulgari SPQMMENFVEKLL-----VAWTEVGLPLQDVSVAACNFCHRPVHFELMSEWERSYFGNMG

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Sciurus_carolin SPQMMENFVEKLL-----VAWTEVGLPLQDVSVAACNFCRHPVHFELMSEWERSYFGNMG
Sciurus_niger@A SPQMMENFVEKLL-----VAWTEVGLPLQDVSVAACNFCRHPVHFELMSEWERSYFGNMG
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Alas2 PQYVTTYAZ
Aplodontia_rufa PQYVTTYA-
Xerus_inauris@A -----
Marmota_monax@A PQYVTTYA-
Spermophilus_da PQYVTTYA-
Ictidomys_tride PQYVTTYA-
Urocitellus_par PQYVTTYA-
Sciurus_vulgari PQYVTTYA-
Sciurus_carolin PQYVTTYA-
Sciurus_niger@A PQYVTTYA-
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CPOX encodes coproporphyrinogen oxidase. Diseases associated with *CPOX* include hereditary coproporphyria and harderoporphyria. Among its related pathways are porphyrin metabolism and biosynthesis of cofactors. Gene ontology (GO) annotations related to this gene include protein homodimerization activity and structural constituent of eye lens. An important paralog of this gene is *ENSG00000285635*. The protein encoded by this gene is the sixth enzyme of the heme biosynthetic pathway. The encoded enzyme is soluble and found in the intermembrane space of mitochondria. This enzyme catalyzes the stepwise oxidative decarboxylation of coproporphyrinogen III to protoporphyrinogen IX, a precursor of heme.

Within a 426-amino acid *CPOX* sequence, we noted eight point-differences among sciurids. All members of *Sciurus* showed addition of a leucine at position 17. The human sequence had an insertion of two amino acids at positions 115-116.

```
Cpox MALRLGRLGSDPWVRA-VLGDYQLRAASPRCASARVCQLPGTAGPQPRRGLGYG-----
Aplodontia_rufa MAWHLGRLSAGSCLRA-ARGGCGELRAWSQRCAGGRICRPPGTAGTESCRGLAHG-----
Xerus_inauris@C MAWHLGRLSVGSCWRA-ARGGCGELRAWSQRCAGGRICRPPGTAGTESCRGLAHG-----
Marmota_monax@C MAWYLGRLSAGPCWRA-ARGGCGELRAWSQRCATGRICQPPGTAGTMHSRGLAHG-----
Spermophilus_da MAWYLGRLSAGPCWRA-ARGGCGELRAWSQRCATGRICQPPGTAGTGHSRGLLEGXXXXX
Ictidomys_tride MAWYLGRLSAGPCWRA-ARGGCGELRAWSQRCATGHICQPPGTAGTVHSRGLAHG-----
Urocitellus_par MAWYLGRLSAGPCWRA-ARGGCGELRAWSQRCATGRICQPPGTAGTVHSRGLAHG-----
Sciurus_vulgari MAWHLGRLSAGPCWRALARGGCGELRAWSQRCAGGRICRPPGTAGTESCRGLAHG-----
Sciurus_carolin MAGHLGRLSAGPCWRALARGGCGELRAWSQRCAGGRICRPPGTAGTESCRGLAHG-----
Sciurus_niger@C MAWHLGRLSAGPCWRALARGGCGELRAWSQRCAGGRICRPPGTAGTESCRGLAHG-----
** ****. .. ** . *. .:*** * *** .:***:****:*. *** *
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Cpox -PWARGGSGLGTRLAATLAGLAGLAAAFAFGHVQRAEMVPKSSGARSPSPGRREEDGDELA
Aplodontia_rufa -PSARGSRWPGTGLTAALAGLAGLATAAVGHVHRAEMVPKSSGARSPSPERPLE--DELA
Xerus_inauris@C -PSAKGGPWPWTGLAAALAGLAGLATAAFGHVQRAEMVPKSSGARSSSQERPLD--DDLA
Marmota_monax@C -PSARGGPWPWTGLAAALAGLAGLATAAFGHVQRAEMVPKSSGARSPSPERPQE--DDLA
Spermophilus_da XXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXAEEMVPKSSGARSPSPVRPQE--DDLA
Ictidomys_tride -PSARGGPWPWTGLAAALAGLAGLATAAFGHVHRAEMVPKSSGARSPSPERPQE--DDLA
Urocitellus_par -PSARGGPWPWTGLAAALAGLAGLATAAFGHVHRAEMVPKSSGARSPSPERPQE--DDLA
Sciurus_vulgari -PSSGGGSRPGTGLATALAGLAGLATAAFGHVQRAEMVPKSSGARSPSPERSQE--DDLT
Sciurus_carolin -PSSGGGSRPGAGLATALAGLAGLATAAFGHVQRAEMVPKSSGVRSPSPERPQE--DDLT
Sciurus_niger@C -PSSGGGSRPGAGLATALAGLAGLATAAFGHVQRAEMVPKSSGVRSPSPERPQE--DDLT
*****.***.* * : *:::
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Cpox RRCSTFMSSPVTELRELRRRPEDMKTKEMLMIMETQAQVCRALAQVDGVADFTVDRWERK
```


Aplodontia_rufa RRCSCFMASPVTDLDELRRKPGDMKTKMELLIILETQAQVCQALAQVDGGANFSVDRWERK
Xerus_inauris@C RRCSCFMASPVTDLSELRRKPGDMKTKMELLIILETQAQVCQALAQVDGGASFSVDRWERK
Marmota_monax@C RRCSCFMASPVTDLDELRRKPGDMKTKMELLIILETQAQVCQALAQVDGVASFSVDRWERK
Spermophilus_da RRCSCFMASPVTDLDELRRKPGDMKTKMELLIILETQAQVCQALAQVDGVASFSVDRWERK
Ictidomys_tride RRCSCFMASPVTDLDELRRKPGDMKTKMELLIILETQAQVCQALAQVDGVASFSVDRWERK
Urocitellus_par RRCSCFMASPVTDLDELRRKPGDMKTKMELLIILETQAQVCQALAQVDGVASFSVDRWERK
Sciurus_vulgari RRCSCFMALPVTDLDELRRKPGDMKTKMELLIILETQAQVCQALAQVDGGASFSVDRWERK
Sciurus_carolin RRCSCFMALPVTDLDELRRKPGDMKTKMELLIILETQAQVCQALAQVDGGASFSVDRWERK
Sciurus_niger@C RRCSCFMALPVTDLDELRRKPGDMKTKMELLIILETQAQVCQALAQVDGGASFSVDRWERK
**** *: **:* **:* **:* **:* **:* **:* **:* **:* **:* **:* **:* **:* **:* **:* **:* **:* **:*

Cpox EGGGGITCVLQDGRVFEKAGVSI SVVHGNSLSEEAANQMRGRGKTLKTKD SKLPFTAMGVS
Aplodontia_rufa EGGGGISCVLQDGRVFEKAGVSI SVVHGNSLSEEAQMRSRGKILKTKDGKLPFTAMGVS
Xerus_inauris@C EGGGGISCVLQDGRVFEKAGVSI SVVHGNSLSEEAQMRSRGKVLKTKNDGKLPFSAMGVS
Marmota_monax@C EGGGGISCVLQDGRVFEKAGVSI SVVHGNSLSEEAQMRSRGKVLKTKDGKLPFCAMGVS
Spermophilus_da EGGGGISCVLQDGRVFEKAGVSI SVVHGNSLSEEAQMRSRGKVLKTKDGKLPFSAMGVS
Ictidomys_tride EGGGGISCVLQDGRVFEKAGVSI SVVHGNSLSEEAQMRSRGKVLKTKDGKLPFSAMGVS
Urocitellus_par EGGGGISCVLQDGRVFEKAGVSI SVVHGNSLSEEAQMRSRGKVLKTKDGKLPFSAMGVS
Sciurus_vulgari EGGGGITCVLQDGRVFEKAGVSI SVVHGNSLSEEAQMRSRGKVLKTKDGKLPFSAMGVS
Sciurus_carolin EGGGGISCVLQDGRVFEKAGVSI SVVHGNSLSEEAQMRSRGKVLKTKDGKLPFSAMGVS
Sciurus_niger@C EGGGGISCVLQDGRVFEKAGVSI SVVHGNSLSEEAQMRSRGKVLKTKDGKLPFSAMGVS
*****:*****:*****:*****:*****:*****:*****:*****:*****:*****:*****

Cpox SVIHPKNPYAPTMHFNYRYFEVEEADGNTHWWFGGGCDLTPTYLNQEDAVHFHRTLKEAC
Aplodontia_rufa SVIHPKNPYAPTMHFNYRYFEVEEADGKQWWFGGGCDLTPTYLNQEDAVHFHRTLKEAC
Xerus_inauris@C SVIHPKNPHAPTIHFNYRYFEVEEADGKQWWFGGGCDLTPTYLNQEDAVHFHRTLKEAC
Marmota_monax@C SVIHPKNPHAPTIHFNYRYFEVEEADGKQWWFGGGCDLTPTYLNQEDAVHFHRTLKEAC
Spermophilus_da SVIHPKNPHAPTIHFNYRYFEVEEADGKQWWFGGGCDLTPTYLNQEDAVHFHRTLKEAC
Ictidomys_tride SVIHPKNPHAPTIHFNYRYFEVEEADGKQWWFGGGCDLTPTYLNQEDAVHFHRTLKEAC
Urocitellus_par SVIHPKNPHAPTIHFNYRYFEVEEADGKQWWFGGGCDLTPTYLNQEDAVHFHRTLKEAC
Sciurus_vulgari SVIHPKNPYAPTMHFNYRYFEVEEADGNQWWFGGGCDLTPTYLNQEDAVHFHRTLKEAC
Sciurus_carolin SVIHPKNPYAPTIHFNYRYFEVEEADGNKQWWFGGGCDLTPTYLNQEDAVHFHRTLKEAC
Sciurus_niger@C SVIHPKNPYAPTIHFNYRYFEVEEADGNKQWWFGGGCDLTPTYLNQEDAVHFHRTLKEAC
*****:*****:*****:*****:*****:*****:*****:*****:*****:*****:*****

Cpox DQHGPDLYPKFKKWCCDYFFI VHRGERRGIGGIFDLDLSPSKEEAFRVKTC AEAVVPS
Aplodontia_rufa DQHGPDLYPKFKKWCCDYFFI VHRGERRGIGGIFDLDLSPSKEEVFRVKS CAQAVVPS
Xerus_inauris@C DQHGPDLYPKFKKWCCDYFFI THRGERRGIGGIFDLDLSPSKEEVFRVQ SCARAVVPS
Marmota_monax@C DQHGPDLYPKFKKWCCDYFFI AHRGERRGIGGIFDLDLSPSKEEVFRVQ SCAQAVVPS
Spermophilus_da DQHGPDLYPKFKKWCCDYFFI AHRGERRGIGGIFDLDLSPSKEEVFRVQ SCAQAVVPS
Ictidomys_tride DQHGPDLYPKFKKWCCDYFFI AHRGERRGIGGIFDLDLSPSKEEVFRVQ SCAQAVVPS
Urocitellus_par DQHGPDLYPKFKKWCCDYFFI AHRGERRGIGGIFDLDLSPSKEEVFRVQ SCAQAVVPS
Sciurus_vulgari DQHGPDLYPKFKKWCCDYFFI THRGERRGIGGIFDLDLSPSKEEAFRVKTC AEAVVPS
Sciurus_carolin DQHGPDLYPKFKKWCCDYFFI THRGERRGIGGIFDLDLSPSKEEAFRVKTC AEAVVPS
Sciurus_niger@C DQHGPDLYPKFKKWCCDYFFI THRGERRGIGGIFDLDLSPSKEEAFRVKTC AEAVVPS
*****:*****:*****:*****:*****:*****:*****:*****:*****:*****:*****

Cpox YVPIVKKHCDDSYTPRDKLWQQLRRGRYVEFNLLYDRGTFKFLFTPGSRIESILMSLPLT
Aplodontia_rufa YIPLVKKHCDDSYTPQEKLWQQLRRGRYVEFNLLYDRGTFKFLFTPGSRIESILMSLPLT
Xerus_inauris@C YIPLVKKHCDDSYTPQEKLWQQLRRGRYVEFNLLYDRGTFKFLFTPGSRIESILMSLPLT
Marmota_monax@C YIPLVKKHCDDSYTPQEKLWQQLRRGRYVEFNLLYDRGTFKFLFTPGSRIESILMSLPLT
Spermophilus_da YIPLVKKHCDDSYTPREKLWQQLRRGRYVEFNLLYDRGTFKFLFTPGSRIESILMSLPLT
Ictidomys_tride YIPLVKKHCDDSYTPQEKLWQQLRRGRYVEFNLLYDRGTFKFLFTPGSRIESILMSLPLT
Urocitellus_par YIPLVKKHCDDSYTPQEKLWQQLRRGRYVEFNLLYDRGTFKFLFTPGSRIESILMSLPLT
Sciurus_vulgari YVPIVKKHCDDSYTPRDKLWQQLRRGRYVEFNLLYDRGTFKFLFTPGSRIESILMSLPLT
Sciurus_carolin YVPLVKKHCDDSYTPQEKLWQQLRRGRYVEFNLLYDRGTFKFLFTPGSRIESILMSLPLT

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Sciurus_niger@C YVPLVKKHCDDSYTPQEKLWQQLRRGRYVEFNLLYDRGTFGLFTPGSRIESILMSLPLT
*:*:*****:*:*:*****
Cpox ARWEYMHSPENSKEAEILEVLRHPKDWVHZ
Aplodontia_rufa AR-----
Xerus_inauris@C AR-----
Marmota_monax@C AR-----
Spermophilus_da AR-----
Ictidomys_tride AR-----
Urocitellus_par AR-----
Sciurus_vulgari ARWEYMHSPENSKAKT--QSLHPPRDWVH-
Sciurus_carolin ARWAYMHLPPENSKEAEILRVLHHPRDWVH-
Sciurus_niger@C ARWAYMHSPENSK-----
**

```

FECH encodes ferrochelatase. Diseases associated with *FECH* include erythropoietic protoporphyria 1 and autosomal erythropoietic protoporphyria. Among its related pathways are porphyrin metabolism and the HIF-1-alpha transcription factor network. Gene ontology (GO) annotations related to this gene include iron ion binding and 2 iron, 2 sulfur cluster binding. The protein encoded by this gene is localized to the mitochondrion, where it catalyzes the insertion of the ferrous form of iron into protoporphyrin IX in the heme synthesis pathway. Two transcript variants encoding different isoforms have been found for this gene.

Among 319 amino acids for FECH, sciurids differed at three. *X. inauris* had a 10-amino acid deletion at positions 256-265, and *M. monax* a 15-amino acid deletion at positions 250-264.

```

Fech MLSASANMAAALRAAGALLREPLVHGSSRACQPWRCQ-SGAAVAATTEKVHHAKTTPKQA
Aplodontia_rufa -----
Xerus_inauris@F -----LVCGNLRASRSWKCKLNTI AVAVDKGKDQCVKNNANPQV
Marmota_monax@F -----LVLGNSKASQSWKCKLNTTAVAVHTGKVQFVKNNKNPQV
Spermophilus_da -----LVLGNSKASQSWKCKLNTTAVAVHTGKVQFVKNNKLNQV
Ictidomys_tride -----
Urocitellus_par -----LVLGNSKASQSWKCKLNTTAVAVHTGKVQFVKNNKNPQV
Sciurus_vulgari -----
Sciurus_carolin -----
Sciurus_niger@F -----

```

```

Fech QPERRKPKTGILMLNMGGPETLGEVQDFLQRLFLDRDLMTLPIONKLAPPIAKRRTPKIQ
Aplodontia_rufa -----SKLAPPIAKRRTPKIQ
Xerus_inauris@F QLEKKKPKGTGI-MLNIGGPETLGEVQDFLQRLFLDRDLIIFPI-NKLAPPIAKRRTPKIQ
Marmota_monax@F QLEKKNPKTGILMLNIGGPETLGEVQDFLQRLFLDRDLMI FPI-NKLAPPIAKRRTPKIQ
Spermophilus_da QLE- KNPKTGILMLNIGGPETLGEVQDFLQRLFLDRDLMI FPI-NKLAPPIAKRRTPKIQ
Ictidomys_tride -----SKLAPPIAKRRTPKIQ
Urocitellus_par QLEKKNPKTGILMLNIGGPETLGEVQDFLQRLFLDRDLMI FPI-NKLAPPIAKRRTPKIQ
Sciurus_vulgari -----SKLAPPIAKRRTPRIQ
Sciurus_carolin -----SKLAPPIAKRRTPRIQ
Sciurus_niger@F -----SKLAPPIAKRRTPRIQ
.*****:**

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```

Fech EQYRRIGGGSPIKMWTSKQEGEMVKLLDELSPATAPHKYYYIGFRYVHPLTEEAIEEMERD
Aplodontia_rufa EQYRRIGGGSPIKMWTSKQEGEMVKLLDDLSPDTAPHKYYYIGFRYVHPLTEEAIDKMERD
Xerus_inauris@F EQYRRIGGGSPIKMWTSKQEGEMVKLLDELSPATAPHKYYYIGFRYVHPLTEEAIEEMERD
Marmota_monax@F EQYRRIGGGSPIKMWTSKQEGEMVKLLDELSPATAPHKYYYIGFRYVHPLTEEAIEEMERD
Spermophilus_da EQYRRIGGGSPIKMWTSKQEGEMVKLLDELSPATAPHKYYYIGFRYVHPLTEEAIEEMERD

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Ictidomys_tride EQYRRIGGGSPIKMWTSKQEGEMVKLLDELSPNTAPHKYYIGFRYVHPLTEEAIEEMERD
Urocitellus_par EQYRRIGGGSPIKMWTSKQEGEMVKLLDELSPATAPHKYYIGFRYVHPLTEEAIEEMERD
Sciurus_vulgari EQYRRIGGGSPIKMWTSKQEGEMVKLLDELSPPTAPHKYYIGFRYVHPLTEEAIDEMERD
Sciurus_carolin EQYRRIGGGSPIKMWTSKQEGEMVKLLDELSPPTAPHKYYIGFRYVHPLTEEAIDEMERD
Sciurus_niger@F EQYRRIGGGSPIKMWTSKQEGEMVKLLDELSPPTAPHKYYIGFRYVHPLTEEAIDEMERD
*****:*** *****:****

Fech GLERAIIFTQYPQYSCSTTGSSLNAIYRYYNEVGKPTMKWSTIDRWPTHPLLIQCFADH
Aplodontia_rufa GLERAIIFTQYPQYSCSTTGSSLNAIYRYYNEVGKPTMKWSTIDRWPTHPLLIQVSDH
Xerus_inauris@F GLERAIIFTQYPQYSCSTTGSSLNAIYRYYNEVGKPTMKWSTIDRWPTHPLLIQCFADN
Marmota_monax@F GLERAIIFTQYPQYSCSTTGSSLNAIYRYYNEVGKPTMKWSTIDRWPTHPLLIQCFADY
Spermophilus_da GLERAIIFTQYPQYSCSTTGSSLNAIYRYYNEVGKPTMKWSTIDRWPTHPLLIQCFADH
Ictidomys_tride GLERAIIFTQYPQYSCSTTGSSLNAIYRYYNEVGKPTMKWSTIDRWPTHPLLIQCFADH
Urocitellus_par GLERAIIFTQYPQYSCSTTGSSLNAIYRYYNEVGKPTMKWSTIDRWPTHPLLIQCFADH
Sciurus_vulgari GLERAIIFTQYPQYSCSTTGSSLNAIYRYYNEVGKPTMKWSTIDRWPTHPLLIQVSDH
Sciurus_carolin GLERAIIFTQYPQYSCSTTGSSLNAIYRYYNEVGKPTMKWSTIDRWPTHPLLIQVSDH
Sciurus_niger@F GLERAIIFTQYPQYSCSTTGSSLNAIYRYYNEVGKPTMKWSTIDRWPTHPLLIQVSDH
*****:***** *

Fech ILKELNHFPEEKRSEVVILFSAHSLPMSVVNRGDPYPQEVGATVHKVMEKLGYPNPYRLV
Aplodontia_rufa ILKELNHFPEEKRSEVVILFSAHSLPLQVNVNRGDPYPQEVGATVHKVMESLGYSNPYRLV
Xerus_inauris@F LQKELHHFPLEKGGH-----QMSVVNRGDPYPQEVGATVHKVMEKLGYPNPYRLV
Marmota_monax@F LQKELHHFP-----LPMSSVVNRGDPYPQEVGATVHKVMERLGYSNPYRLV
Spermophilus_da ILKELNHFPEEKRSEVVILFSAHSLPMSVVNRGDPYPQEVGATVHKVMERLGYSNPYRLV
Ictidomys_tride ILKELNHFPEEKRSEVVILFSAHSLPMSVVNRGDPYPQEVGATVHKVMERLGYSNPYRLV
Urocitellus_par ILKELNHFPEEKRSEVVILFSAHSLPMSVVNRGDPYPQEVGATVHKVMERLGYSNPYRLV
Sciurus_vulgari ILKELNHFPEEKRSEVVILFSAHSLPMQVVNRGDPYPQEVGATVHKVMERLGYSNPYRLV
Sciurus_carolin ILKELNHFPEEKRSEVVILFSAHSLPMQVVNRGDPYPQEVGATVHKVMERLGYSNPYRLV
Sciurus_niger@F ILKELNHFPEEKRSEVVILFSAHSLPMQVVNRGDPYPQEVGATVHKVMERLGYSNPYRLV
: ***:*** :.***** *****.*****

Fech WQSKVGPVWLGQPQTDEAIKGLCERGRKNILLVPIAFTSDHIETLYELDI EYSQVLAQKC
Aplodontia_rufa WQSKVGPVSWLGPQTDEAIKGLCERGRKNILLVPIAFTSDHIETLYELDI EYSQILAKEC
Xerus_inauris@F WQSKVGPVWLGQPQTDEAIKGLCERGRKNILLVPIAFTSDHIETLYELDI EYSQVLAQKC
Marmota_monax@F WQSKVGPVWLGQPQTDEAIKGLCERGRKNILLVPIAFTSDHIETLYELDI EYSQVLAQKY
Spermophilus_da WQSKVGPVWLGQPQTDEAIKGLCERGRKNILLVPIAFTSDHIETLYELDI EYSQVLAKEV
Ictidomys_tride WQSKVGPVWLGQPQTDEAIKGLCERGRKNILLVPIAFTSDHIETLYELDI EYSQVLAKEV
Urocitellus_par WQSKVGPVWLGQPQTDEAIKGLCERGRKNILLVPIAFTSDHIETLYELDI EYSQVLAKEC
Sciurus_vulgari WQSKVGPVWLGQPQTNEAIKGLCERGRKNILLVPIAFTSDHIETLYELDI EYSQVLAKEV
Sciurus_carolin WQSKVGPVWLGQPQTNEAIKGLCERGRKNILLVPIAFTSDHIETLYELDI EYSQVLAKEC
Sciurus_niger@F WQSKVGPVWLGQPQTNEAIKGLCERGRKNILLVPIAFTSDHIETLYELDI EYSQVLAKEC
*****.*****:*****:***:.

Fech GAENIRRAESLNGNPLFSKALADLVHSHIQSNKLCSTQLSLNCLCVNPVCRKTKSFFTS
Aplodontia_rufa GAENIRRAESLNGNPLFSQALADLVHSHIQSNRLCSKQLTLCSCPLCVNPVCRETKSFFTN
Xerus_inauris@F GAENIMKRESFAGNPLFSQALADLVSHIQSNKLCSTQLTLCSCPLCVNPVCRETKSFFTN
Marmota_monax@F -VENNKRGESLNGNPLFSQALADLVLSHIQSNKLCSTQLTLCSCPLCVNPVCRETKSFFTN
Spermophilus_da GAENIRRAESLNGNPLFSQALADLVLSHIQSNKLCSTQLTLCSCPLCVNPVCRETKSFFTN
Ictidomys_tride GAENIRRAESLNGNPLFSQALADLVLSHIQSNKLCSTQLTLCSCPLCVNPVCRETKSFFTN
Urocitellus_par GAENIRRAESLNGNPLFSQALADLVLSHIQSNKLCSTQLTLCSCPLCVNPVCRETKSFFTN
Sciurus_vulgari STENIRRAESLNGNPLFSQALADLVHSHIQSNTQCSKQLTLCSCPLCVNPVCRETKSFFTN
Sciurus_carolin GAENIRRAESLNGNPLFSQALADLVHSHIQSNTLCSKQLTLCSCPLCVNPVCRETKSFFTN
Sciurus_niger@F GAENIRRAESLNGNPLFSQALADLVHSHIQSNTLCSKQLTLCSCPLCVNPVCRETKSFFTN
.** : **: *****:***** ***** **.***:*****:*****.

Fech QQLZ

Aplodontia_rufa QQL-
 Xerus_inauris@F QQL-
 Marmota_monax@F QQL-
 Spermophilus_da QQL-
 Ictidomys_tride QQL-
 Urocitellus_par QQL-
 Sciurus_vulgari QQM-
 Sciurus_carolin QQM-
 Sciurus_niger@F QQM-
 **:

HMBS encodes hydroxymethylbilane synthase. Diseases associated with *HMBS* include acute intermittent porphyria and mood disorder. Among its related pathways are porphyrin metabolism and biosynthesis of cofactors. Gene ontology (GO) annotations related to this gene include hydroxymethylbilane synthase activity. The encoded protein is the third enzyme of the heme biosynthetic pathway and catalyzes the head-to-tail condensation of four porphobilinogen molecules into the linear hydroxymethylbilane. Alternatively spliced transcript variants encoding different isoforms have been described.

Among 242 amino acids in *Sciurus niger* *HMBS*, there were two point-differences among *Sciurus* species and an insertion of 37 amino acids in *S. vulgaris*. There were major amino acid sequence insertions among all species, including 37 amino acids in *A. rufa*; five amino acids for *M. monax*, *I. dauricus*, and *U. parryi*; and eight amino acids for *S. dauricus*.

```
Hmbs          MSGNGGAATTAEENGSKMRVIRVGTRKSQLARIQTDTVVAMLKALYPGIQFEIIAMSTTG
Aplodontia_rufa -----
Xerus_inauris@H -----
Marmota_monax@H -----
Spermophilus_da -----
Ictidomys_tride -----
Urocitellus_par -----
Sciurus_vulgari -----
Sciurus_carolin -----
Sciurus_niger@H -----
```

```
Hmbs          DKILDOTALSKIGEKSLFTKELENALEKNEVDLVVHSLKDVPTILPPGFTIGAICKRENPC
Aplodontia_rufa -----ICPSFHDLPSFPRRENPY
Xerus_inauris@H -----RRENPC
Marmota_monax@H -----RRENPC
Spermophilus_da -----RRENPC
Ictidomys_tride -----RRENPC
Urocitellus_par -----RRENPC
Sciurus_vulgari -----
Sciurus_carolin -----RRENPC
Sciurus_niger@H -----RRENPC
```

```
Hmbs          DAVVFHPKFIGKTLETLEPEK-----SAV
Aplodontia_rufa DAVVFHPKFIGKTLETLEPEKRVGPEWAAWWDVHRRWKAGRSRKGTVTICISFTFSLSSSTV
Xerus_inauris@H DAVVFHPKFIGKTLETLEPEK-----SAV
Marmota_monax@H DAVVFHPKFIGKTLETLEPEK-----SAV
Spermophilus_da DAVVFHPKFIGKTLETLEPEK-----SAV
```

```
Ictidomys_tride DAVVFHPKFVVGKTLETLEPEK-----SAV
Urocitellus_par DAVVFHPKFVVGKTLETLEPEK-----SAV
Sciurus_vulgari -----
Sciurus_carolin DAVVFHPKFVVGKTLETLEPEK-----SAV
Sciurus_niger@H DAVVFHPKFVVGKTLETLEPEK-----SAV
```

```
Hmbs GTSSLRRAAQLQRKFPHPLEFKSI-----RGNLNTRLRKLDELQEFSAIIVLAVAGLQRMGW
Aplodontia_rufa GTSSLRRAAQLQRKFPHPLEFKNI-----RGNLNTRLRKLDEQHEFSAIILAVAGLQRMGW
Xerus_inauris@H GTSSLRRAAQLQRKFPHPLEFKSI-----RGNLNTRLRKLDEQHEFSAIILAVAGLQRMGW
Marmota_monax@H GTSSLRRAAQLQRKFPHPLEFKSIVSFRRRGNLNTRLRKLDEQHEFSAIILAVAGLQRMGW
Spermophilus_da GTSSLRRAAQLQRKFPHPLEFKSIVSFRRRGNLNTRLRKLDEQHEFSAIILAVAGLQRMGW
Ictidomys_tride GTSSLRRAAQLQRKFPHPLEFKSI-----RGNLNTRLRKLDEQHEFSAIILAVAGLQRMGW
Urocitellus_par GTSSLRRAAQLQRKFPHPLEFKSIVSFRRRGNLNTRLRKLDEQHEFSAIILAVAGLQRMGW
Sciurus_vulgari -----RGNLNTRLRKLDEQHEFSAIILAVAGLQRMGW
Sciurus_carolin GTSSLRRAAQLQRKFPHPLEFKSI-----RGNLNTRLRKLDEQHEFSAIILAVAGLQRMGW
Sciurus_niger@H GTSSLRRAAQLQRKFPHPLEFKSI-----RGNLNTRLRKLDEQHEFSAITLAVAGLQRMGW
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```

```
Hmbs QNRVGQI-----LHPEECMYAVGQGALAVEVRAKDQDILDVGVVLDHPETLLRC
Aplodontia_rufa QNRVGQVGASPLPSPIYLLPPPCFFSVGQGALGVEVRAKDQDILDVGVVLDHPETLLRC
Xerus_inauris@H QNRVGQV-----GAPNECMYAVGQGALGVEVRAKDQDILDVGVVLDHPETLLRC
Marmota_monax@H QNRVGQV-----LHPEECMYAVGQGALAVEVRAKDQDILDVGVVLDHPETLLRC
Spermophilus_da QNRVGQV-----LHPEECMYAVGQGALAVEVRAKDQDILDVGVVLDHPETLLRC
Ictidomys_tride QNRVGQI-----VERRSSLMCFHQGALAVEVRAKDQDILDVGVVLDHPETLLRC
Urocitellus_par QNRVGQI-----LERRSSLMCFHQGALAVEVRAKDQDILDVGVVLDHPETLLRC
Sciurus_vulgari QNRVGQV-----LHPEECMYAVGQGALGVEVRAKDQDILDVGVVLDHPETLLRC
Sciurus_carolin QNRVGQV-----LHPEECMYAVGQGALGVEVRAKDQDILDVGVVLDHPETLLRC
Sciurus_niger@H QNRVGQV-----LHPEECMYAVGQGALGVEVRAKDQDILDVGVVLDHPETLLRC
*****: .: .. *****.*****.*****
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```
Hmbs IAERAFLRHL-----EGGCSVPVAVHTVMKDG-----
Aplodontia_rufa IAERAFLRHL-----EGGCSVPVAVHTVMKDG-----
Xerus_inauris@H IAERAFLRHL-----EGGCSVPVAVHTVMKDG-----
Marmota_monax@H IAERAFLRHL-----EGGCSVPVAVHTVIKDG-----
Spermophilus_da IAERAFLRHL-----EGGCSVPVAVHTVIKDG-----
Ictidomys_tride IAERAFLRHLVGPVVLVLEEGGCSVPVAVHTVIKDG-----
Urocitellus_par IAERAFLRHL-----EGGCSVPVAVHTVIKDG-----
Sciurus_vulgari IAERAFLRHL-----EGGCSVPVAVHTVMKDGQVSSGKWAGRQRTGISPYASQVSKQ
Sciurus_carolin IAERAFLRHL-----VGGCSVPVAVHTVMKDG-----
Sciurus_niger@H IAERAFLRHL-----VGGCSVPVAVHTVMKDG-----
***** *****:***
```

```
Hmbs -----QLYLTGGVWSLDGSDSMQETMQATIQVPVQQEDGPEDDPQLVGITARN
Aplodontia_rufa -----QLYLTGGVWSLDGSDSMQETMQATIHVPVQHEGPEDDPQLVGITARN
Xerus_inauris@H -----QLYLTGGVWSLDGSDSMQETMQATINVPVQYEDGPEDDPQLVGITARN
Marmota_monax@H -----QLYLTGGVWSLDGSDSMQETMQATIHVPVQYDHGPEDDPQLVGITARN
Spermophilus_da -----QLYLTGGVWSLDGSDSMQETMQATIHVPVQYDDGPEDDPQLVGITARN
Ictidomys_tride -----QLYLTGGVWSLDGSDSMQETMQATIHVPTQQYDGPEDDPQLVGITARN
Urocitellus_par -----QLYLTGGVWSLDGSESMQETMQATIHVPVQYDNGPEDDPQLVGITARN
Sciurus_vulgari SPLRICNNYFLS QLYLTGGVWSLDGSDSMQETMQATIHVPVQYEDGPEDDPQLVGITARN
Sciurus_carolin -----QLYLTGGVWSLDGSDSMQETMQATIQVPIQYEDGPEDDPQLVGITARN
Sciurus_niger@H -----QLYLTGGVWSLDGSDSMQETMQATIQVPIQYEDGPEDDPQLVGITARN
*****:*** * .***** *
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```
Hmbs IPRGAQLAAENLGISLASLLLNKGAKNILDVARQLNDVRZ
```

```

Aplodontia_rufa IPRGAQLAAENLGISLANLLLLNKGAKNILDVARQLNDVH-
Xerus_inauris@H IPRGAQLAAENLGISLANLLLLNKGAKNILDVARQLNDV--
Marmota_monax@H IPRGAQLAAENLGISLASLLLLNKGAKNILDVARQLNDVH-
Spermophilus_da IPRGAQLAAENLGISLASLLLLNKGAKNILDVARQLNDVH-
Ictidomys_tride IPRGAQLAAENLGISLASLLLLNKGAKNILDVARQLNDVH-
Urocitellus_par IPRGAQLAAENLGISLASLLLLNKGAKNILDVARQVNDVH-
Sciurus_vulgari IPRGAQLAAENLGISLANLLLLNKGAKNILDVARQLNDV--
Sciurus_carolin IPRGAQLAAENLGISLANLLLLNKGAKSILDVARQLNDV--
Sciurus_niger@H IPRGAQLAAENLGISLANLLLLNKGAKSILDVARQLNDV--
*****.*****.*****:***
    
```

PPOX encodes protoporphyrinogen oxidase. Among its related pathways are porphyrin and biosynthesis of cofactors. Gene ontology (GO) annotations related to this gene include oxidoreductase activity and oxygen-dependent protoporphyrinogen oxidase activity. This gene encodes the penultimate enzyme of heme biosynthesis, which catalyzes the 6-electron oxidation of protoporphyrinogen IX to form protoporphyrin IX. Mutations in this gene cause variegate porphyria, an autosomal dominant disorder of heme metabolism resulting from a deficiency in protoporphyrinogen oxidase, an enzyme located on the inner mitochondrial membrane. Alternatively spliced transcript variants encoding the same protein have been identified.

Relative to a 557-amino acid sequence for *S. niger*, we observed four point-differences among sciurids. There were insertions of 31 amino acids for *M. monax* and *X. inauris*, 25 for *S. niger*, 24 for *M. monax* and *I. tridecemlineatus*, and 25 for *X. inauris*. There was a deletion of 32 amino acids for *A. rufa*; of 35 for *A. rufa*, *X. inauris*, and *I. tridecemlineatus*; of eight amino acids in *A. rufa*, *X. inauris*, and *I. tridecemlineatus*; and of 26 amino acids for *M. monax* and *S. dauricus*.

```

Ppox          MGRTVIVLGGGISGLAASYHLIRGSPSPKVVILVEGSKRLGGWIRSIRGSDGAI FELGPRG
Aplodontia_rufa -----SPPCRQVVLVEGSKRLGGWIRSVRGPDGAIFELGPRG
Xerus_inauris@P -----PSCRQVVLVEGSKRLGGWIRSVRGPDGAIFELGPRG
Marmota_monax@P -----QVVLVEGSKRLGGWIRSIRGPGGAI FELGPRG
Spermophilus_da -----QVVLVEGSKRLGGWIRSIRGPGGAI FELGPRG
Ictidomys_tride -----QVVLVEGSKRLGGWIRSIRGPGGAI FELGPRG
Urocitellus_par -----QVVLVEGSKRLGGWIRSIRGPGGAI FELGPRG
Sciurus_vulgari -----PSCRQVVLVEGSKRLGGWIRSVRGPDGAIFELGPRG
Sciurus_carolin -----AASDGTTMPASCRQVVLVEGSKRLGGWIRSVRGPDGAIFELGPRG
Sciurus_niger@P -----PSCRQVVLVEGSKRLGGWIRSVRGPDGAIFELGPRG
: * : ***** : * * . *****
    
```

```

Ppox          IRPAGALGARTLLLSELGLESEVLPVRGDHPAAQNRFLYVGGTLHPLPSGLRGLLRPSP
Aplodontia_rufa IRPAGALGARTLLLSELGLESEVLPVRGDHPAAQNRFLYVGGTLHPLPSGLRGLLRPSP
Xerus_inauris@P IRPAGPLGARTLLLSELGLESEVLPVRGDHPAAQNRFLYVGGTLHPLPSGLRGLLRPSP
Marmota_monax@P IRPAGVLGARTLLLSELGLESEVLPVRGDHPAAQNRFLYVGGTLHPLPSGLRGLLRPSP
Spermophilus_da IRPAGVLGARTLLLSELGLESEVLPVRGDHPAAQNRFLYVGGTLHPLPSGLRGLLRPSP
Ictidomys_tride IRPAGVLGARTLLLSELGLESEVLPVRGDHPAAQNRFLYVGGTLHPLPSGLRGLLRPSP
Urocitellus_par IRPAGVLGARTLLLSELGLESEVLPVRGDHPAAQNRFLYVGGTLHPLPSGLRGLLRPSP
Sciurus_vulgari IRPAGALGARTLLLSELGLESEVLPVRGDHPAAQNRFLYVGGTLHPLPSGLRGLLRPSP
Sciurus_carolin IRPAGALGARTLLLSELGLESEVLPVRGDHPAAQNRFLYVGGTLHPLPSGLRGLLRPSP
Sciurus_niger@P IRPAGALGARTLLLSELGLESEVLPVRGDHPAAQNRFLYVGGTLHPLPSGLRGLLRPSP
*****
    
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```

Ppox          PFSKPLFWAGLRELLKPRGKEPDET VHSFAQRRLGPEVASLAMDSL CRGVFAGNSRELSI
Aplodontia_rufa PFSKSLWTGLRELLKPRGKEPDET VHSFAQRRLGPEVASLVMDSL CRGVFAGNSRELSV
    
```

Xerus_inauris@P PFSKPLLWAGLRELTKPRGKEPDETVHSFAQRRLGPEVASLAMDSLRCRVFAGNSRELSI
 Marmota_monax@P PFSKPLLWAGLRELTKPRGKEPDETVHSFAQRRLGPEVASLAMDSLRCRVFAGNSRELSI
 Spermophilus_da PFSKPLLWAGLRELTKPRGKEPDETVHSFAQRRLGPEVASLAMDSLRCRVFAGNSRELSI
 Ictidomys_tride PFSKPLLWAGLRELTKPRGKEPDETVHSFAHRRLGPEVASLVMDSLRCRVFAGNSRELSI
 Urocitellus_par PFSKPLLWAGLRELTKPRGKEPDETVHSFAQRRLGPEVASLVMDSLRCRVFAGNSRELSI
 Sciurus_vulgari PFSRPLLWAGLRELTKPRSKEPDETVHSFAQRRLGPEVASLAMDSLRCRVFAGNSRELSI
 Sciurus_carolin PFSRPLLWAGLRELTKPRSKEPDETVHSFAQRRLGPEVASLAMDSLRCRVFAGNSRELSI
 Sciurus_niger@P PFSRPLLWAGLRELTKPRSKEPDETVHSFAQRRLGPEVASLAMDSLRCRVFAGNSRELSI
 :.*:.*:** ***.*****:*****.***:*****:

Ppox RSCFPSLFQAEQTHRSILLGLLLG-----AGQSP
 Aplodontia_rufa RSCFPSLFQAEQTHRSILLGLLLG-----AGQSP
 Xerus_inauris@P RSCFPSLFQAEQTYRSIILGLLLGAGEEGLVQRVKILRTARVRQGVLRFFFVAVSLGQSP
 Marmota_monax@P RSCFPSLFQAEQTHRSVILGLLFS-----SGQSP
 Spermophilus_da RSCFPSLFQAEQTHRSVILGLLFS-----SGKSP
 Ictidomys_tride RSCFPSLFQAEQTHRSVILGLLFS-----SGKSP
 Urocitellus_par RSCFPSLFQAEQTHRSVILGLLFS-----SGKSP
 Sciurus_vulgari RSCFPSLFQAEQTHRSIILGLLLG-----AGQSP
 Sciurus_carolin RSCFPSLFQAEQTHRSIILGLLLG-----AGQSP
 Sciurus_niger@P RSCFPSLFQAEQTHRSIILGLLLGAGE-----VAGRYGLPVRQGVLIFFFAAASAGQSP
 *****:.*:.*:***** ***.*****:*****.***:*****:

Ppox QPDSALIRQARAERWSQWSLRGGLEVLQPALHNHLASKGVTVLSGQPVCGLSLQPEGRWK
 Aplodontia_rufa QPDSALVRQARAERWSQWSLRGGMEMLPQTLHTYLT SRGVSVLKGQPVCGLTQAEGHWK
 Xerus_inauris@P QSDSALIRQAQAERWSQWSLRGGMEMLPQTLDAYLTSRGVSVLRGQPVCGLTQAEGRWK
 Marmota_monax@P QPDSALIRQARAERWSQWSLRGGMETLPQTLDTYLT SRGVSVLRGQPVCGLSLQAEHGWK
 Spermophilus_da QPDSALIRQARAERWSQWSLRGGMETLPQTLDTYLT SRGVSVLRGQPVCGLSLQAEHGWK
 Ictidomys_tride QPDSALIRQARAERWSQWSLRGGMETLPQTLDTYLT SRGVSVLRGQPVCGLSLQAEHGWK
 Urocitellus_par QPDSALIRQARAERWSQWSLRGGMETLPQTLDTYLT SRGVSVLRGQPVCGLSLQAEHGWK
 Sciurus_vulgari QPDSALIRQARAERWSQWSLRGGMEMLPQTLDTYLT SNGVSVLRGQPVCGLTQAEGHWK
 Sciurus_carolin QPDSALIRQARAERWSQWSLRGGMEMLPQTLDTYLT SNGVSVLRGQPVSGLTQAEGHWK
 Sciurus_niger@P QPDSALIRQARAERWSQWSLRGGMEMLPQTLDTYLT SNGVSVLRGQPVSGLTQAEGHWK
 *.***:.*:.*:***** ***.*****:*****.***:*****:

Ppox VSLGDSSEADHIIISAI PASELSKLLPAEAAPLARILSTIKAVSVAVVNLQYRGACLPV-
 Aplodontia_rufa VSLGDSSEADHIIISAI PASELSKLLPAEAAPLAHILSTITAVSVAVVNLQYRGARLPV-
 Xerus_inauris@P VSLGDSSEADHIIISAI PASELSKLLPMEAAPLAQVLSTITAVSVAVVNLQYQGARLPVQ
 Marmota_monax@P VSLGDSSEADHIIISAI PASELSKLLPTEAAPLAQVLNTITAVSVAVVNLQYRGARLPVQ
 Spermophilus_da VSLGDSSEADHIIISAI PASELSKLLPTEAAPLAQVLNTITAVSVAVVNLQYRGARLPV-
 Ictidomys_tride VSLGDSSEADHIIISAI PASELSKLLPTEAAPLAQVLNNTITAVSVAVVNLQYRGACLPVQ
 Urocitellus_par VSLGDSSEADHIIISAI PASELSKLLPTEAAPLAQVLNTITAVSVAVVNLQYRGARLPV-
 Sciurus_vulgari VSLGDSSEADHIIISAI PASELSKLLPAEAAPLARILSTIKAVSVAVVNLQYRGACLP-
 Sciurus_carolin VSLGDSSEADHIIISAI PASELSKLLPAEAAPLARILSTIKAVSVAVVNLQYRGACLP-
 Sciurus_niger@P VSLGDSSEADHIIISAI PASELSKLLPAEAAPLARILSTIKAVSVAVVNLQYRGACLP-
 *****:.*:.*:***** ***.*****:*****.***:*****:

Ppox -----QGFGHLVPSSEDPVTLGIVYDSVAFPEQDGNPPSLR
 Aplodontia_rufa -----QGFGHLVPSSEDPGVLGIVYDSVAFPEQDGSSPGLR
 Xerus_inauris@P VYWRGKEGIDSDGTFPSFFFFPPLQGFGHLVPSSEDPGVLGIVYDSVAFPEQDGSPGLR
 Marmota_monax@P VYRGEKKVPQMEFFLLI-FFFFPPLQGFGHLVPSSEDPGVLGIVYDSVAFPEQDGSPGLR
 Spermophilus_da -----QGFGHLVPSSEDPGVLGIVYDSVAFPEQDGSPGLR
 Ictidomys_tride VYGGGKKGVASDGVFPS-YFFFSLQGFGHLVPSSEDPGVLGIVYDSVAFPEQDGSSPGLR
 Urocitellus_par -----QGFGHLVPSSEDPGVLGIVYDSVAFPEQDGSPGLR
 Sciurus_vulgari -----QGFGHLVPSSEDPGVLGIVYDSVAFPEQDGSPGLR
 Sciurus_carolin -----QGFGHLVPSSEDPGVLGIVYDSVAFPEQDGNPPGLR
 Sciurus_niger@P -----QGFGHLVPSSEDPGVLGIVYDSVAFPEQDGNPPGLR

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Ppox VT-----VMLGGYWLQKLKAAGHQLSPELFQQQ
Aplodontia_rufa VTVRA-----GGYWLQKLKAAGHQLSPELFQQQ
Xerus_inauris@P VTVRAGHVWVAFQRRALLCPVVGKAKLVSAIF-APVMLGGSWLQMLEASGCVLSQELFQQQ
Marmota_monax@P VTVRAGHVWVAFQRRALLCSIVGKAKLVSAIFPSQVMLGGSWLQMLEASGCVLSQELFQKQ
Spermophilus_da VTVRAGHFVWVAFQRRALFCPIVGGKAKLVSAIFPSQVMLGGSWLQMLEASGYVLSRELFQQQ
Ictidomys_tride VTVRAGHVWVAFQRTLFCPIVGGKAKLVSAIFPSQVMLGGSWLQMLEASGCVLSQELFQQQ
Urocyonellus_par VTVRAGHVWVAFQRRALFCPIVGGKAKLVSAIFPSQVMLGGSWLQMLEASGCVLSQELFQQQ
Sciurus_vulgari VTVRAGHVVRVVFQRRAPLCPVGGKAKLVSAIFPSQVMLGGSWLQMLEASGCVLSQKLFQQQ
Sciurus_carolin VTVRAGHVVRVVFQRRAPLCPVGGKAKLVSAIFPPQVMLGGSWLQMLEASDCVLSQKLFQQQ
Sciurus_niger@P VTVRAGHVVRVVFQRRAPLCPVGGKAKLVSAIFPPQVMLGGSWLQMLEASGCVLSQKLFQQQ
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Ppox AQEAAATQLGLKEPPSHCLVHLHK-----
Aplodontia_rufa AQEAAATQLGLKEPPSHCLVHLHQ-----
Xerus_inauris@P AQEAVATQLGLKEPPSHCLVHLHK-----
Marmota_monax@P AQEAVATQLGLKEPPSHCLVHLHKVSWGKFLSVPCCRPCIQGIFVTVQCQPRPGKSIINFI
Spermophilus_da AQEAVATQLGLKEPPSHCLVHLHKVSWGKFLSVPYCRPCIQGIFVTVQCQPRPGKSIINFI
Ictidomys_tride AQEAVATQLGLKEPPSHCLVHLHK-----
Urocyonellus_par AQEAVATQLGLKEPPSHCLVHLHKVSWGKFLSVPYCRPCIQGIFVTVQCQPRPGKSIINFI
Sciurus_vulgari AQEAVATQLGLKEPPSHCLVHLHKVSWGKFLSVPFLRP-IQGIFVILC-PRPGKSVINFI
Sciurus_carolin AQEAVATQLGLKEPPSHCLVHLHKVSWGKFLSVPFWRP-IQGIFVTVQC-PRPGKSVINFI
Sciurus_niger@P AQEAVATQLGLKEPPSHCLVHLHKVSWGKFLSVPFWRP-IQGIFVTVQC-PRPGKSVINFI
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Ppox -----NCIPQYTIHGWQK-----LDSAMQFLTAQRL
Aplodontia_rufa -----NCIPQYTLGHWQKLGKLGKLGKGGQRKSDL-KLHSFHPFLPESAMKFLSAQKL
Xerus_inauris@P -----NCIPQYTLGHWQKLGKLGKQLGELRKSDDLQPQNSFHPFLLESAMKFLAAQRL
Marmota_monax@P LYPLSSQNCIPQYTLGHWQK-----LESAMKFLAAQRL
Spermophilus_da LYPLSSQNCIPQYTLGHWQK-----LESAMKFLAAQRL
Ictidomys_tride -----NCIPQYTLGHWQKLGKVGKQLGGLRK-DL-PQNSFYFSLLESAMKFLAAQRL
Urocyonellus_par LYPLSSQNCIPQYTLGHWQKLGKVGKQLGGLRK-DL-PQNSFYFSLLESAMKFLGAQRL
Sciurus_vulgari LYSPLSSQNCIPQYTLGHWQKLGKFGKQLGGLKKSDDL-PQNSFYFPLPESAMKFLAAQKL
Sciurus_carolin LYPLSSQNCIPQYTLGHWQKLGKFGKQLGGLKKSDDL-PQNSFHPFLPESAMKFLAAQKL
Sciurus_niger@P LYPLSSQNCIPQYTLGHWQKLGKFGKQLGGLKKSDDL-PQNSFHPFLPESAMKFLAAQKL
*****:***** :***:** **:
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Ppox PLTLGASYEGVAVNDCIESGRQAAVAVLGTESNSZ
Aplodontia_rufa PLTLGASYEGVAVNDCIESGRQAAVSVLGTESNS-
Xerus_inauris@P PLTLGASYEGVAVNDCIESGRQAAISVLGPESNS-
Marmota_monax@P PLTLGASYEGVAVNDCIESGRQAAISVLGPESNS-
Spermophilus_da PLTLGASYEGVAVNDCIESGRQAAISVLGPESNS-
Ictidomys_tride PLTLGASYEGVAVNDCIESGRQAAISVLGPESNS-
Urocyonellus_par PLTLGASYEGVAVNDCIESGRQAAISVLGPESNS-
Sciurus_vulgari PLTLGASYEGVAVNDCIESGRQAAISVLGPESN--
Sciurus_carolin PLTLGASYEGVAVNDCIESGRQAAISVLGPES---
Sciurus_niger@P PLTLGASYEGVAVNDCIESGRQAAISVLGPESN--
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UROD encodes uroporphyrinogen decarboxylase. Diseases associated with *UROD* include porphyria cutanea tarda and familial porphyria cutanea tarda. Among its related pathways are biosynthesis of cofactors and metabolism. Gene ontology (GO) annotations related to this gene include uroporphyrinogen decarboxylase activity. This gene encodes an enzyme responsible for catalyzing the conversion of uroporphyrinogen to coproporphyrinogen through the removal of

four carboxymethyl side chains. Mutations and deficiency in this enzyme are known to cause familial porphyria cutanea tarda and hepatoerythropoetic porphyria.

Within a 426-amino acid *UROD* sequence for *S. niger*, we observed nine point-differences among sciurids, including one contiguous tract of six where the *S. niger* sequence differed from those of *S. vulgaris* and *S. carolinensis*. *S. carolinensis* had a deletion of 38 amino acids relative to the sequences of *S. vulgaris* and *S. niger*. Comparing among all species, we observed insertions of 15 and 15 amino acids in *A. rufa* and 50 in *S. dauricus* and *U. parryii*.

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Urod      MEANGFGLQNFPELKNNTFLRAAWGEETDYTPVWCMRQAGRYL-----P-----
Aplodontia_rufa  -----QTFPELKNNTFLRAAWGEETDYTPVWCMRQAGRYLPGQGQGPEVINVCV----
Xerus_inauris@U -----QSFPELKNNTFLRAAWGEETDYTPVWCMRQAGRYLPGQGQGP-----
Marmota_monax@U -----QNFPELKNNTFLRAAWGEETDYTPVWCMRQAGRYL-----P-----
Spermophilus_da -----QNFPELKNNTFLRAAWGEETDYTPVWCMRQAGRYL-----P-----
Ictidomys_tride -----QNFPELKNNTFLRAAWGEETDYTPVWCMRQAGRYL-----P-----
Urocitellus_par -----QNFPELKNNTFLRAAWGEETDYTPVWCMRQAGRYL-----P-----
Sciurus_vulgari -----QSFPELKNNTFLRAAWGEETDYTPVWCMRQAGRYLPGEGQGPEFITLGSGGW
Sciurus_carolin -----QSFPELKNNTFLRAAWGEETDYTPVWCMRQAGRYLPGQGQGPEFITLGSGGW
Sciurus_niger@U -----QSFPELKNNTFLRAAWGEETDYTPVWCMRQAGRYLPGQGQGPEFITLGSGGW
                * .*****:*****
    
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Urod      -----EFRETRAAQDFSTCRSPEACCELT-----
Aplodontia_rufa  -MSRASRLRSVFLFLFAEFKTRATQDFSTCRSPEACCELT-----
Xerus_inauris@U -----EFRETRAAQDFSTCRSPEACCELTQVRGPEREVFMPSAYRLP
Marmota_monax@U -----EFRETRATQDFSTCRSPEACCELTQVRGPPQREVFMPTACHLA
Spermophilus_da -----EFRETRATQDFSTCRSPEACCELTQVRSPQREVFMPSACHLA
Ictidomys_tride -----EFRETRATQDFSTCRSPEACCELTQVRGPPQREVFMPSACHLA
Urocitellus_par -----EFRETRATQDFSTCRSPEACCELTQVRGPPQREVFMPSACHLA
Sciurus_vulgari  GGALGARLRS AFLLSAEFRETATQDFSTCRSPEACCELTQVRGPPQREVFMSSACHLA
Sciurus_carolin  GGALGARLRS AFLLSAEFRETATQDFSTCRSPEACCELT-----
Sciurus_niger@U  GGALGARLRS AFLLSAEFRETATQDFSTCRSPEACCELT-----
                ***:***:*****
    
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```

Urod      -----LQPLRRFPLDAAIIFSDILVVPQALGMEVTMVPKGKPSFPEPLREERDLERLRDP
Aplodontia_rufa  -----LQPLRRFPLDAAIIFSDILVVPQALGLEVTMEPGKGKPSFPEPLREERDLERLRDP
Xerus_inauris@U  TCFHSLQPLRRFPLDAAIIFSDILVVPQALGMEVIMVPKGKPSFPEPLREERDLERLRDP
Marmota_monax@U  TRFHSLQPLRRFPLDAAIIFSDILVVPQALGMEVTMVPKGKPSFPEPLREERDLERLRDP
Spermophilus_da  TRFHSLQPLRRFPLDAAIIFSDILVVPQALGMEVTMVPKGKPSFPEPLREERDLERLRDP
Ictidomys_tride  TRFHSLQPLRRFPLDAAIIFSDILVVPQALGMEVTMVPKGKPSFPEPLREERDLERLRDP
Urocitellus_par  TRFHSLQPLRRFPLDAAIIFSDILVVPQALGMEVTMVPKGKPSFPEPLREERDLERLRDP
Sciurus_vulgari  TCFHSLQPLRRFPLDAAIIFSDILVVPQALGMEVTMVPKGKPSFSEPLREEQDLERLRDP
Sciurus_carolin  -----LQPLRRFPLDAAIIFSDILVVPQALGMEVTMVPKGKPSFSEPLREERDLERLRDP
Sciurus_niger@U  -----LQPLRRFPLDAAIIFSDILVVPQALGMEVTMVPKGKPSFSEPLREERDLERLRDP
                *****:*** * *****:*****
    
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Urod      AAAASELG YVFQAITLTRQQLAGRVPLIGFAGAP-----
Aplodontia_rufa  AAVASELNYVFQAITLTRQQLAGRVPLIGFAGAPVIWEEQGLGHEITLADLVLTREGGVS
Xerus_inauris@U  SVVASELGYVFQAITLTRQQLAGRVPLIGFAGAP-----
Marmota_monax@U  EVVTSELGYVFQAITLTRQQLAGRVPLIGFAGAP-----
Spermophilus_da  EVVTSELGYVFQAITLTRQQLAGRVPLIGFAGAPVIWEKQGIGALWHI--WSRGRKCQCG
Ictidomys_tride  EVVTSELGYVFQAITLTRQQLAGRVPLIGFAGAPVIWEKQGTGALWHI--WCRGRKCECG
Urocitellus_par  EVVTSELGYVFQAITLTRQQLAGRVPLIGFAGAPVTWEKQGIGALWHI--WCRGRKCQCG
Sciurus_vulgari  AVVASELGYVFQAITLTRQQLAGRVPLIGFAGAPVIWDKQGTGTDHAG--ISSIDKGSVN
Sciurus_carolin  AVVASELGYVFQAITLTRQQLAGRVPLIGFAGAP-----
Sciurus_niger@U  AVVASELGYVFQAITLTRQQLAGRVPLIGFAGAPVTWDKQGTGADHAG--VSSIDKGSVS
                ..:***.*****:*****
    
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```

Urod -----WTLMTYMVEGGSSSTMAQAKRWLYQRPQASHKLLGILTDVLPVPLYIGQ
Aplodontia_rufa LTSALHLYILL-WTLMTYMVEGGSSSTMAQSKRWLYQRPQASHKLLGILTDALVPYLVGQ
Xerus_inauris@U -----WTLMTYMVEGGSSSTMAQSKRWLYQRPQASHQLLRILTALVPYLVGQ
Marmota_monax@U -----WTLMTYMVEGGSSSTMAQSKRWLYQRPQASHQLLRILTALVPYLVGQ
Spermophilus_da FCTVTSL-ILLQWTLMTYMVEGGSSSTMAQSKRWLYQRPQASHQLLRILTALVPYLVGQ
Ictidomys_tride FCTVTSL-ILLQWTLMTYMVEGGSSSTMAQSKRWLYQRPQASHQLLRILTALVPYLVGQ
Uroditellus_par FCTVTSL-ILLQWTLMTYMVEGGSSSTMAQSKRWLYQRPQASHQLLRILTALVPYLVGQ
Sciurus_vulgari VASALRLCILLQWTLMTYMVEGGSSSTMAQSKRWLYQRPQASHQLLRILTALVPYLI--
Sciurus_carolin -----WTLMTYMVEGGSSSTMAQSKRWLYQRPQASHQLLRILTALVPYLI--
Sciurus_niger@U VASALHLCILLQWTLMTYMVEGGSSSTMAQSKRWLYQRPQASHQLLRILTALVPYLVGQ
*****.*****:*****:*** ***.*****:
    
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Urod VAAGA-QALQLFESHAGHLGTELFKSFALPYIRDVAKRVKAGLQKAGLAPVPM-----
Aplodontia_rufa VPRQVPRHCSSFESHAGHLGTELFKSFALPYIRDVAKRVKAGLQKAGLAPVPM-----
Xerus_inauris@U VAAGA-QALQLFESHAGHLGQPLFNKFSALPYIRDVAKRVKARLQKAGLAPVPM-----
Marmota_monax@U VAAGA-QALQLFESHAGHLGQPLFNKFSALPYIRDVAKRVKARLQKAGLAPVPM-----
Spermophilus_da VAAGA-QALQLFESHAGHLGQPLFNKFSALPYIRDVAKRVKARLQKAGLAPVPMVRIGIGW
Ictidomys_tride VAAGA-QALQLFESHAGHLGQPLFNKFSALPYIRDVAKRVKARLQKAGLAPVPM-----
Uroditellus_par VAAGA-QALQLFESHAGHLGQPLFNKFSALPYIRDVAKRVKARLQKAGLAPVPMVRIGIGW
Sciurus_vulgari FFTTL-LALQLFESHAGHLGQPLFSKFSALPYIRDVAKRVKAKLQKAGLAPVPM-----
Sciurus_carolin FFTTL-LALQLFESHAGHLGQPLFSKFSALPYIRDVAKRVKAKLQKAGLAPVPM-----
Sciurus_niger@U VAAGA-QALQLFESHAGHLGQPLFSKFSALPYIRDVAKRVKAKLQKAGLAPVPM-----
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Urod -----IIFAKDGHFALEELAQ
Aplodontia_rufa -----IIFAKDGHFALEELAQ
Xerus_inauris@U -----IIFAKDGHFALEELAQ
Marmota_monax@U -----IIFAKDGHFALEELAQ
Spermophilus_da WVCRVFRVSLAWTEVTTGGQQKCIQERLVGTARPYVSQIRFFLQIIIFAKDGHFALEELAQ
Ictidomys_tride -----IIFAKDGHFALEELAQ
Uroditellus_par WVCRVFRVSHAWTEVTTGGQQKCIQERLVGTARPYVSQILFFLQIIIFAKDGHFALEELAQ
Sciurus_vulgari -----IIFAKDGHFALEELAQ
Sciurus_carolin -----IIFAKDGHFALEELAQ
Sciurus_niger@U -----IIFAKDGHFALEELAQ
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Urod AGYEVVGLDWTVAPKKARERVGKAVTLQGNLDPCALYASEEEEIGRLVQQLDDFGPQRYI
Aplodontia_rufa AGYEVVGLDWTVAPKKARERVGKAVTLQGNLDPCALYASEEEEIGRLVKQMLDDFGPQHYI
Xerus_inauris@U AGYEVVGLDWTMAPKKARERVGKAVTLQGNLDPCALYASEEEEIGRLVQQLDDFGPQRYI
Marmota_monax@U AGYEVVGLDWTVAPKKARERVGKSVTLQGNLDPCALYASEEEEIGRLVQQLDDFGPKRYI
Spermophilus_da AGYEVVGLDWTVAPKKARERVGKSVTLQGNLDPCALYASEEEEIGRLVQQLDDFGPKRYI
Ictidomys_tride AGYEVVGLDWTVAPKKARERVGKSVTLQGNLDPCALYASEEEEIGRLVQQLDDFGPKRYI
Uroditellus_par AGYEVVGLDWTVAPKKARERVGKSVTLQGNLDPCALYASEEEEIGQLVQQLDDFGPKRYI
Sciurus_vulgari AGYEVVGLDWTVAPKKARERVGKAVTLQGNLDPCALYASEEEEIGQMVQKMLDDFGPQRYI
Sciurus_carolin AGYEVVGLDWTVAPKKARERVGKAVTLQGNLDPCALYASEEEEIGQMVQKMLDDFGPQRYI
Sciurus_niger@U AGYEVVGLDWTVAPKKARERVGKAVTLQGNLDPCALYASEEEEIGQMVQKMLDDFGPQRYI
*****:***** *****:*****:***:*****:***
    
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```

Urod ANLGHGLYPDMDPERVGFVDAVHKHSRLLRQNZ
Aplodontia_rufa ANLGHGLYPDMDPEHVGFVDAVHRHSRLLRQN-
Xerus_inauris@U ANLGHGLYPDMDPEHVGFVDAVHRHSRLLRQN-
Marmota_monax@U ANLGHGLYPDMDPEHVGFVDAVHRHSRLLRQN-
Spermophilus_da ANLGHGLYPDMDPEHVGFVDAVHRHSRLLRQN-
Ictidomys_tride ANLGHGLYPDMDPEHVGFVDAVHRHSRLLRQN-
Uroditellus_par ANLGHGLYPDMDPEHVGFVDAVHRHSRLLRQN-
    
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Sciurus_vulgari ANLGHGLYPDMDPEHVGA FVDAVHRHSRLLRQN-
Sciurus_carolin ANLGHGLYPDMDPEHVGA FVDAVHRHSRLLRQN-
Sciurus_niger@U ANLGHGLYPDMDPEHVGA FVDAVHRHSRLLRQN-
                  *****:*****:*****
    
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UROS encodes uroporphyrinogen III synthase. Diseases associated with *UROS* include porphyria and congenital erythropoietic and cutaneous porphyria, or Gunther’s disease. Among its related pathways are porphyrin metabolism and biosynthesis of cofactors. Gene ontology (GO) annotations related to this gene include cofactor binding and uroporphyrinogen-III synthase activity. The protein encoded by this gene catalyzes the fourth step of porphyrin biosynthesis in the heme biosynthetic pathway. The human *UROS* protein has 199 amino acids before any squirrel one. *M. monax*, *S. dauricus*, *I. tridecemlineatus*, and *U. parryii* had an 11-amino acid sequence not similar to the human sequence. From amino acid 419, the sequences of all species were similar. The *S. niger* amino acid sequence differed from those of *S. vulgaris* and *S. carolinensis* at amino acid 229, and the *S. carolinensis* sequence from those of *S. vulgaris* and *S. niger* at amino acid 256.

```

Uros                MKVLLLLKDAKEDDSGLDPYIQELRLCGLEATLIPVLSFEFMSLPSLSEKLSHPEGFGGLI
Aplodontia_rufa    -----
Xerus_inauris@U    -----
Marmota_monax@U    -----
Spermophilus_da    -----
Ictidomys_tride    -----
Urocyon_vulpes     -----
Sciurus_vulgari    -----
Sciurus_carolin    -----
Sciurus_niger@U    -----
    
```

```

Uros                FTSPRAVEAVKLCLEKDNKTEAWEKSLKDRWNAKSVYVVGSAATSLVNKIGLDAEGAGSG
Aplodontia_rufa    -----
Xerus_inauris@U    -----
Marmota_monax@U    -----
Spermophilus_da    -----
Ictidomys_tride    -----
Urocyon_vulpes     -----
Sciurus_vulgari    -----
Sciurus_carolin    -----
Sciurus_niger@U    -----
    
```

```

Uros                NAEKLAEYICSKPSELPLLFP CGTIKGD TLPKMLKDKGIPMESMHVYQTVPHPGIQGSL
Aplodontia_rufa    -----
Xerus_inauris@U    -----
Marmota_monax@U    -----
Spermophilus_da    -----
Ictidomys_tride    -----
Urocyon_vulpes     -----
Sciurus_vulgari    -----
Sciurus_carolin    -----
Sciurus_niger@U    -----
    
```

```

Uros          KSYIEDQGIPASITFFSPSGLKYSLEYIQALSGSSFDQIKFIAIGPSTTRAMAAKGLPVS
Aplodontia_rufa -----QFAAIGPSTARALAAKGLPVS
Xerus_inauris@U -----SSLPSLQFAAIGPTTARALTSKGLPVS
Marmota_monax@U -----GVMYRKALAEA---LSLPSLQFAAIGPTTAHALTAKGLPVS
Spermophilus_da -----GVMYRKALAEA---LSLPSLQFAAIGPTTAHALTAKGLPVS
Ictidomys_tride -----GVMYQKALAEA---LSLPSLQFAAIGPTTAHALTAKGLPVS
Urocitellus_par -----GVTYRKALAEA---LSLPSLQFAAIGPTTAHALTAKGLPVS
Sciurus_vulgari -----SSLPSSQFAAIGPTTARALAAQGLPVS
Sciurus_carolin -----FAAIGPTTARALAAQGLPVS
Sciurus_niger@U -----SLSSSQFAAIGPTTARTLAAQGLPVS
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Uros          CTAESPTPQALAAGIRNVLPKPNHCCZ
Aplodontia_rufa CTAESPTPQALAASIRKALQPHSCC-
Xerus_inauris@U CTAESPTPQALAAGIRKALQPQGCC-
Marmota_monax@U CTAESPTPQALAAGIRKALQPQGCC-
Spermophilus_da CTAESPTPQALAAGIRKALQPQGCC-
Ictidomys_tride CTAESPTPQALAAGIRKALQPQGCC-
Urocitellus_par CTAESPTPQALAAGIRKALQPQGCC-
Sciurus_vulgari CTAESPTPQALAAGIRKALQPQ----
Sciurus_carolin CTAESPTPQALAAGIWKALQPQGSC-
Sciurus_niger@U CTAESPTPQALAAGIRKALQPQGSC-
*****~::~:~::~:
    
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